

Analisi e Modificazioni di Sequenze di DNA o di RNA con la

TEORIA TRICROMATICA DELL'EQUILIBRIO DEI SISTEMI



Capitolo I ° (Parte Prima):

***Analisi della Sequenza n°1/1
della Catena A dell'Insulina***

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INTRODUZIONE

Lo scopo di questo lavoro è quello di presentare un'importante applicazione della **TEORIA TRICROMATICA DELL'EQUILIBRIO DEI SISTEMI (T.T.E.S.)** all'**analisi e alle modificazioni di sequenze di DNA o di RNA.**

Con il software della **T.T.E.S.** è possibile analizzare una sequenza di DNA (o di RNA) in *maniera innovativa.*

A partire dalla **sequenza originaria** di DNA o di RNA, il software della **T.T.E.S.** genera numerose e diverse **nuove sequenze** di DNA o di RNA **che rispettano fedelmente i diversi e numerosi "trend non manifesti" della sequenza originaria.**

Questo risultato è possibile per **due motivi:**

- 1) perché ogni specifica sequenza di DNA (o di RNA) può essere "trasformata" in numerose e diverse nuove sequenze, rispettando i diversi e numerosi "trend non manifesti" della specifica sequenza originaria (Fig. 1);
- 2) perché ogni specifico "trend non manifesto" della sequenza originaria può generare numerose e diverse nuove sequenze (Fig. 1).

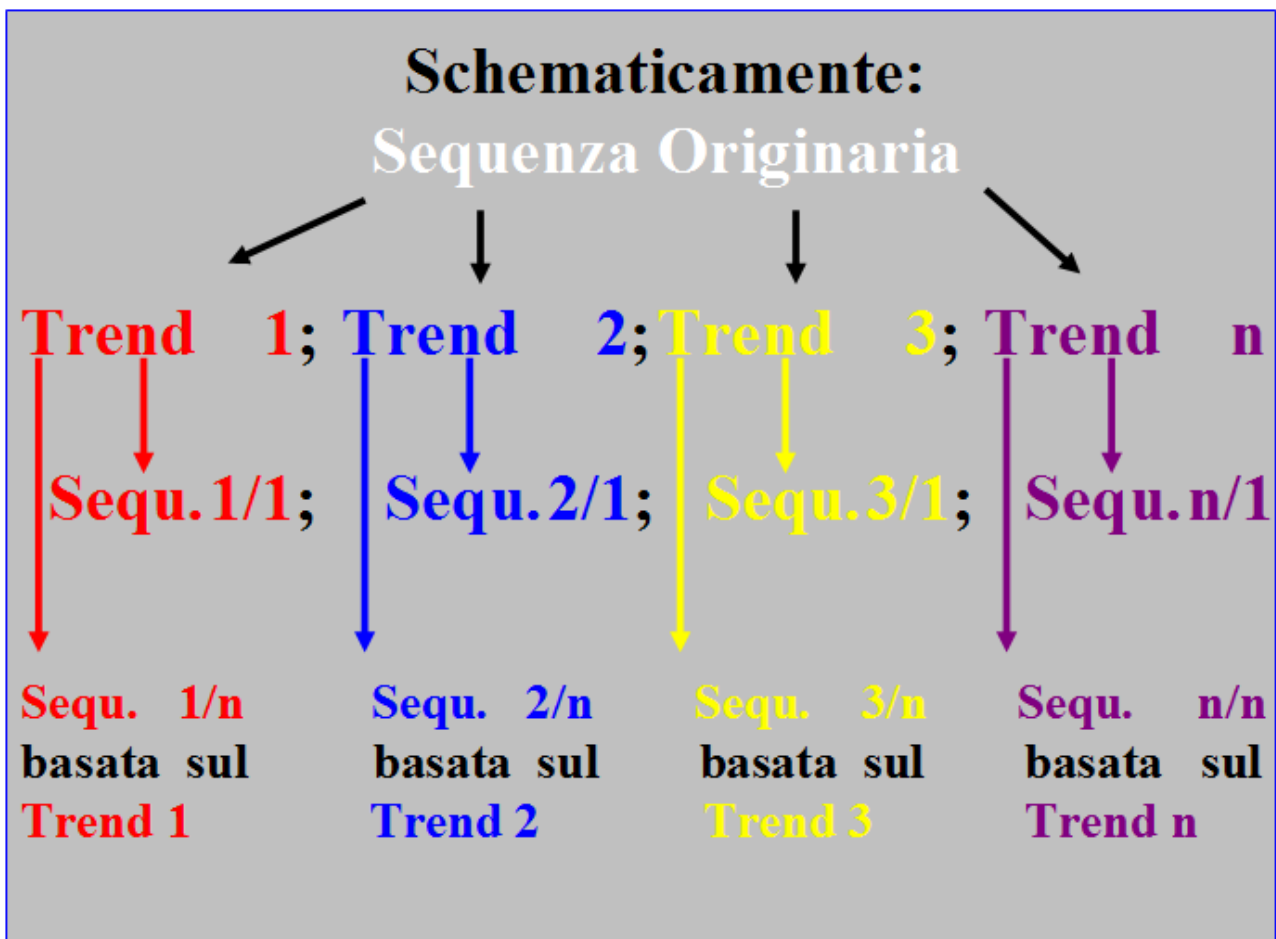


Fig. 1

Queste **nuove sequenze** di DNA (o di RNA) sono costituite da basi quasi totalmente differenti.

L'**ipotesi** che s'intende avvalorare, in questo e negli altri capitoli che seguiranno, è quella che **le nuove sequenze** (generate rispettando fedelmente ognuno dei possibili specifici "trend non manifesti" della sequenza originaria) **abbiano forti relazioni con le caratteristiche della sequenza originaria**.

Queste *nuove sequenze di DNA* (o di RNA) possono, *parzialmente o totalmente*, essere utilizzate per scopi di **ricerca scientifica, industriale, alimentare, ecc.**

Nel Capitolo I° di questo lavoro, *che costituisce solo il primo di molti altri capitoli che seguiranno*, delle numerose e diverse **nuove sequenze generate**, si è deciso di analizzarne solo una tra le possibili: la **Sequenza n°1/1**.

I risultati della ricerca **BLAST** (*Basic Local Alignment Search Tool* (1)) sulla **Sequenza n°1/1** hanno evidenziato allineamenti significativi con il DNA (o RNA) di diversi *organismi*.

La **ricerca bibliografica** ha confermato l'esistenza d'**importanti relazioni** tra le caratteristiche di due *organismi* "esempio" (*Pseudomonas* e *Heligmosomoides polygyrus*) identificati con la ricerca Blast eseguita sulla **Sequenza n°1/1** e alcune caratteristiche funzionali dell'**Insulina**.

In conclusione, l'analisi (attraverso la **T.T.E.S.**) della *sequenza originaria* basata su uno dei suoi "trend non manifesti" (*il Trend n°1*), la creazione di una *nuova sequenza* di DNA (la **Sequenza n°1/1**) dal **Trend n°1** della *sequenza originaria* e la *congruenza* con i dati ottenuti dall'*approfondimento bibliografico* aprono prospettive totalmente inesplorate riguardo all'ambito della ricerca genetica e delle sue innumerevoli applicazioni.

(1) Altschul S. F., Madden T. L., Schaffer A. A., Zhang J., Zhang Z., Miller W. and D. J. Lipman. Gapped BLAST and PSI-BLAST: a new generation of protein database search programs. *Nucleic Acids Res.*, 1997, 25 (17) :3389-3402. PMID: 9254694. <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC146917/>

CAPITOLO I °

(Parte Prima)

1.1 LA TEORIA TRICROMATICA DELL'EQUILIBRIO DEI SISTEMI (T.T.E.S.)

La **T.T.E.S.** è una teoria dei sistemi con la quale è possibile osservare, analizzare, controllare e modificare lo stato di qualsiasi sistema (<http://www.ttesystems.eu/>).

La **T.T.E.S.** è stata applicata, per la prima volta, all'analisi del Sistema Nervoso Vegetativo attraverso l'ausilio del biofeedback periferico

(Il Biofeedback Periferico e la Teoria Tricromatica dell'Equilibrio del Sistema Nervoso Vegetativo; Il Futuro del Biofeedback Periferico: La Teoria Tricromatica dell'Equilibrio del Sistema Nervoso Vegetativo;

L'iperventilazione: un modello privilegiato per la valutazione quantitativa e qualitativa dell'attivazione psicofisiologica con la Teoria Tricromatica dell'Equilibrio del Sistema Nervoso Vegetativo;).

Della **T.T.E.S.** sono previste molte altre applicazioni (<http://www.ttesystems.eu/applicazioni.php>) e alcune di esse sono in corso di sperimentazione e futura pubblicazione.

Per calcolare e per rappresentare visivamente tutte le variazioni di un sistema, la **T.T.E.S.** si è servita del modello di colore **RGB**.

Il modello **RGB** è un metodo per definire i colori basato su *TRE Colori Primari* (Rosso, Verde e Blu).

Il **CUBO** è il solido usato per rappresentare visivamente tutte le possibili variazioni dello stato di un sistema (Fig. 2).

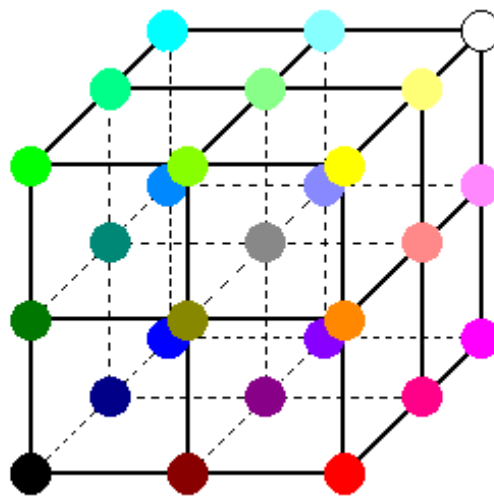


Fig. 2

Per descrivere sinteticamente tutti i possibili stati funzionali di un sistema sono stati utilizzati **8 CODICI PRINCIPALI** (Fig.3) dei **64 CODICI TOTALI**.

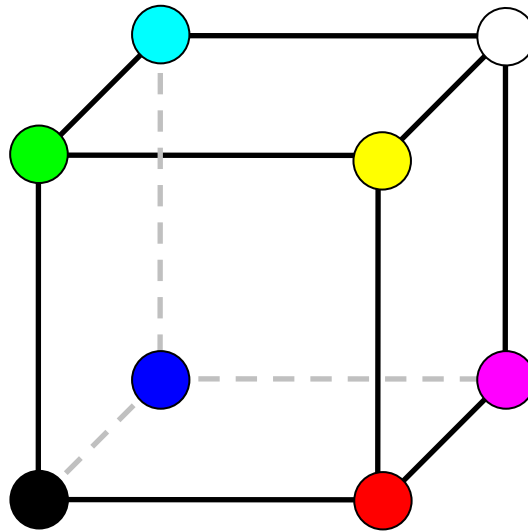


Fig. 3 *

Nei paragrafi successivi è presentato un esempio di elaborazione di una sequenza di DNA con la **T.T.E.S.**.

* Tratto e modificato da: <https://commons.wikimedia.org/wiki/File:Avl3119color4a.jpg>

1.2 ESEMPIO DI ELABORAZIONE DI UNA SEQUENZA DI DNA CON LA T.T.E.S.

L'**acquisizione della sequenza di DNA (o di RNA)** da analizzare o da modificare rappresenta la *prima fase* del processo di analisi con la **T.T.E.S.**

L'acquisizione della sequenza può essere effettuata direttamente dal sito del **NCBI** (*National Center for Biotechnology Information* (2)) o da qualsiasi altra fonte.

Supponiamo di aver deciso di analizzare la seguente **sequenza di 63 basi**:

ggcatcgtggagcagtgctgcaccagcatctgttcctctaccagctggagaactactgcaac

Le **63 basi, tutte codificanti (CDS<1..63)**, dell'intera sequenza corrispondono alla ben nota **Catena A dell'Insulina**.

L'**insulina** è formata da *due catene* polipeptidiche unite da *due ponti disolfuro*:

la **catena A** di **21** aminoacidi (**GIVEQCCTSICSLYQLENYCN**)

e

la **catena B** di **30** aminoacidi (**FVNQHLCGSHLVEALYLVCGERGFFYTKPT**).

L'**insulina** è un *ormone* anabolico di natura *proteica* secreto dalle *cellule beta* delle *isole di Langerhans* del pancreas.

Tra le sue attività principali, l'*insulina* usa il *glucosio* per produrre *energia*, riduce la *glicemia*, blocca la *glicogenolisi*, favorisce la *glicogenosintesi* e stimola la *proliferazione cellulare*.

La **sequenza di 63 basi** di DNA della **Catena A dell'Insulina** è stata oggetto di ricerca **BLAST**.

(2) National Center for Biotechnology Information (**NCBI**)[Internet]. Bethesda (MD): National Library of Medicine (US), National Center for Biotechnology Information; [1988]. Available from: <https://www.ncbi.nlm.nih.gov/>

Parametri della ricerca BLAST:

Programme	<i>Blastn</i>
Word size	<i>11</i>
Expect value	<i>10</i>
Hitlist size	<i>100</i>
Match/Mismatch scores	<i>2,-3</i>
Gapcosts	<i>5,2</i>
Low Complexity Filter	<i>Yes</i>
Filter string	<i>L;m;</i>
Genetic Code	<i>1</i>

Risultati della ricerca BLAST:

Query = **ggcatcgtggagcagtgctgcaccagcatctgttcctctaccagctggagaactactgcaac**

Length = 63

Sequences producing significant alignments:	Score	E
	(Bits)	Value
XM_021081278.1 PREDICTED: Sus scrofa insulin (INS), transcrip...	114	1e-22
https://www.ncbi.nlm.nih.gov/nuccore/1191854326/		
NM_001109772.1 Sus scrofa insulin (INS), mRNA	114	1e-22
https://www.ncbi.nlm.nih.gov/nuccore/172073147/		
AY242112.1 Sus scrofa EWB tyrosine hydroxylase (TH) gene, exo...	114	1e-22
https://www.ncbi.nlm.nih.gov/nuccore/33242556/		
AY242111.1 Sus scrofa H205 tyrosine hydroxylase (TH) gene, ex...	114	1e-22
https://www.ncbi.nlm.nih.gov/nuccore/33242553/		
AY242110.1 Sus scrofa H254 tyrosine hydroxylase (TH) gene, ex...	114	1e-22
https://www.ncbi.nlm.nih.gov/nuccore/33242550/		
AY242108.1 Sus scrofa LRJ tyrosine hydroxylase (TH) gene, exo...	114	1e-22
https://www.ncbi.nlm.nih.gov/nuccore/33242543/		
AY242107.1 Sus scrofa LW1224 tyrosine hydroxylase (TH) gene, ...	114	1e-22
https://www.ncbi.nlm.nih.gov/nuccore/33242539/		
AY242106.1 Sus scrofa LW1461 tyrosine hydroxylase (TH) gene, ...	114	1e-22
https://www.ncbi.nlm.nih.gov/nuccore/33242535/		
AY242105.1 Sus scrofa LW197 tyrosine hydroxylase (TH) gene, e...	114	1e-22
https://www.ncbi.nlm.nih.gov/nuccore/33242531/		
AY242104.1 Sus scrofa LW209 tyrosine hydroxylase (TH) gene, e...	114	1e-22
https://www.ncbi.nlm.nih.gov/nuccore/33242527/		
AY242103.1 Sus scrofa LW3 tyrosine hydroxylase (TH) gene, exo...	114	1e-22
https://www.ncbi.nlm.nih.gov/nuccore/33242523/		

AY242102.1	Sus scrofa LW33361 tyrosine hydroxylase (TH) gene,...	114	1e-22
	https://www.ncbi.nlm.nih.gov/nuccore/33242519/		
AY242101.1	Sus scrofa LW419 tyrosine hydroxylase (TH) gene, e...	114	1e-22
	https://www.ncbi.nlm.nih.gov/nuccore/33242515/		
AY242100.1	Sus scrofa LW463 tyrosine hydroxylase (TH) gene, e...	114	1e-22
	https://www.ncbi.nlm.nih.gov/nuccore/33242511/		
AY242099.1	Sus scrofa M220 tyrosine hydroxylase (TH) gene, ex...	114	1e-22
	https://www.ncbi.nlm.nih.gov/nuccore/33242508/		
AY242098.1	Sus scrofa P208 tyrosine hydroxylase (TH) gene, ex...	114	1e-22
	https://www.ncbi.nlm.nih.gov/nuccore/33242504/		
AY044828.1	Sus scrofa tyrosine hydroxylase gene, partial cds;...	114	1e-22
	https://www.ncbi.nlm.nih.gov/nuccore/21956486/		
AF064555.1	Sus scrofa preproinsulin mRNA, partial cds	114	1e-22
	https://www.ncbi.nlm.nih.gov/nuccore/3885493/		
XM_003909376.4	PREDICTED: Papio anubis insulin (INS), transcr...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/1220171442/		
XM_009185350.3	PREDICTED: Papio anubis insulin (INS), transcr...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/1220171441/		
XM_017948138.2	PREDICTED: Papio anubis insulin (INS), transcr...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/1220171440/		
XM_020883287.1	PREDICTED: Odocoileus virginianus texanus insu...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/1187570232/		
XM_020883286.1	PREDICTED: Odocoileus virginianus texanus insu...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/1187570230/		
XM_015503336.1	PREDICTED: Marmota marmota marmota insulin (LO...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/984133697/		
XM_015503335.1	PREDICTED: Marmota marmota marmota insulin (LO...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/984133695/		
XM_015434180.1	PREDICTED: Macaca fascicularis insulin (INS), ...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/982292203/		
XM_015113354.1	PREDICTED: Macaca mulatta insulin (INS), mRNA	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/966983385/		
XM_011721319.1	PREDICTED: Macaca nemestrina insulin (INS), tr...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795558861/		
XM_011721318.1	PREDICTED: Macaca nemestrina insulin (INS), tr...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795558858/		
XM_011721317.1	PREDICTED: Macaca nemestrina insulin (INS), tr...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795558851/		
XM_011721316.1	PREDICTED: Macaca nemestrina insulin (INS), tr...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795558842/		
XM_012041172.1	PREDICTED: Cercocebus atys insulin (INS), tran...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795499572/		
XM_012041171.1	PREDICTED: Cercocebus atys insulin (INS), tran...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795499567/		
XM_012041169.1	PREDICTED: Cercocebus atys insulin (INS), tran...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795499562/		

XM_011930076.1	PREDICTED: Colobus angolensis palliatus insuli...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795357485/		
XM_011930075.1	PREDICTED: Colobus angolensis palliatus insuli...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795357479/		
XM_011930074.1	PREDICTED: Colobus angolensis palliatus insuli...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795357473/		
XM_011988228.1	PREDICTED: Mandrillus leucophaeus insulin (INS...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795176057/		
XM_011988227.1	PREDICTED: Mandrillus leucophaeus insulin (INS...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795176052/		
XM_011988226.1	PREDICTED: Mandrillus leucophaeus insulin (INS...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795176047/		
XM_011988225.1	PREDICTED: Mandrillus leucophaeus insulin (INS...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/795176041/		
XM_008004634.1	PREDICTED: Chlorocebus sabaeus insulin (INS), ...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/635011293/		
XM_008004561.1	PREDICTED: Chlorocebus sabaeus insulin (INS), ...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/635011291/		
XM_006141067.1	PREDICTED: Tupaia chinensis insulin (INS), mRNA	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/562822569/		
XM_004278069.1	PREDICTED: Orcinus orca insulin (INS), mRNA	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/466049859/		
AY242109.1	Sus scrofa JWB tyrosine hydroxylase (TH) gene, exo...	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/33242547/		
NM_001282255.1	Ictidomys tridecemlineatus insulin (Ins), mRNA	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/532691821/		
NM_001284919.1	Macaca fascicularis insulin (INS), mRNA	109	5e-21
	https://www.ncbi.nlm.nih.gov/nuccore/548960960/		
XM_022591057.1	PREDICTED: Delphinapterus leucas insulin (INS)...	105	6e-20
	https://www.ncbi.nlm.nih.gov/nuccore/1246201225/		
XM_022591055.1	PREDICTED: Delphinapterus leucas insulin (INS)...	105	6e-20
	https://www.ncbi.nlm.nih.gov/nuccore/1246201223/		
XM_022591054.1	PREDICTED: Delphinapterus leucas insulin (INS)...	105	6e-20
	https://www.ncbi.nlm.nih.gov/nuccore/1246201221/		
NM_001130093.2	Canis lupus familiaris insulin (INS), mRNA	105	6e-20
	https://www.ncbi.nlm.nih.gov/nuccore/1212980167/		
XM_021152514.1	PREDICTED: Mus caroli insulin-1 (LOC110286053)...	105	6e-20
	https://www.ncbi.nlm.nih.gov/nuccore/1195703068/		
XM_017887811.1	PREDICTED: Rhinopithecus bieti insulin (INS), ...	105	6e-20
	https://www.ncbi.nlm.nih.gov/nuccore/1059087102/		
XM_017887804.1	PREDICTED: Rhinopithecus bieti insulin (INS), ...	105	6e-20
	https://www.ncbi.nlm.nih.gov/nuccore/1059087100/		
XM_017887801.1	PREDICTED: Rhinopithecus bieti insulin (INS), ...	105	6e-20
	https://www.ncbi.nlm.nih.gov/nuccore/1059087098/		
XM_017887795.1	PREDICTED: Rhinopithecus bieti insulin (INS), ...	105	6e-20
	https://www.ncbi.nlm.nih.gov/nuccore/1059087095/		

XM_017887787.1 PREDICTED: Rhinopithecus bieti insulin (INS), ... https://www.ncbi.nlm.nih.gov/nuccore/1059087093/	105	6e-20
XM_017669266.1 PREDICTED: Manis javanica insulin (INS), mRNA https://www.ncbi.nlm.nih.gov/nuccore/1048405902/	105	6e-20
XM_004759671.2 PREDICTED: Mustela putorius furo insulin (INS)... https://www.ncbi.nlm.nih.gov/nuccore/859820716/	105	6e-20
XM_013060037.1 PREDICTED: Mustela putorius furo insulin (INS)... https://www.ncbi.nlm.nih.gov/nuccore/859820712/	105	6e-20
XM_010382053.1 PREDICTED: Rhinopithecus roxellana insulin (IN... https://www.ncbi.nlm.nih.gov/nuccore/724917366/	105	6e-20
XM_007171156.1 PREDICTED: Balaenoptera acutorostrata scammoni... https://www.ncbi.nlm.nih.gov/nuccore/594635595/	105	6e-20
AC188659.9 Canis familiaris chromosome 18, clone XX-484I11, c... https://www.ncbi.nlm.nih.gov/nuccore/126032460/	105	6e-20
AC187029.8 Canis familiaris chromosome 18, clone XX-127C6, co... https://www.ncbi.nlm.nih.gov/nuccore/148245384/	105	6e-20
DQ250565.1 Mus caroli preproinsulin 1 (Ins1) gene, complete cds https://www.ncbi.nlm.nih.gov/nuccore/82749721/	105	6e-20
X61092.1 C.aethiops gene for preproinsulin https://www.ncbi.nlm.nih.gov/nuccore/X61092.1	105	6e-20
V00179.1 Dog gene encoding insulin https://www.ncbi.nlm.nih.gov/nuccore/V00179.1	105	6e-20
XM_014795833.1 PREDICTED: Ceratotherium simum simum insulin (... https://www.ncbi.nlm.nih.gov/nuccore/XM_014795833.1	104	2e-19
XM_022507720.1 PREDICTED: Enhydra lutris kenyoni insulin (LOC... https://www.ncbi.nlm.nih.gov/nuccore/XM_022507720.1	100	3e-18
XM_021685179.1 PREDICTED: Neomonachus schauinslandi insulin (... https://www.ncbi.nlm.nih.gov/nuccore/XM_021685179.1	100	3e-18
XM_021215010.1 PREDICTED: Mus pahari insulin-1 (LOC110333420)... https://www.ncbi.nlm.nih.gov/nuccore/XM_021215010.1	100	3e-18
XM_019036301.1 PREDICTED: Gorilla gorilla gorilla insulin (IN... https://www.ncbi.nlm.nih.gov/nuccore/XM_019036301.1	100	3e-18
XM_019036300.1 PREDICTED: Gorilla gorilla gorilla insulin (IN... https://www.ncbi.nlm.nih.gov/nuccore/XM_019036300.1	100	3e-18
XM_004050427.2 PREDICTED: Gorilla gorilla gorilla insulin (IN... https://www.ncbi.nlm.nih.gov/nuccore/XM_004050427.2	100	3e-18
XM_004050428.2 PREDICTED: Gorilla gorilla gorilla insulin (IN... https://www.ncbi.nlm.nih.gov/nuccore/XM_004050428.2	100	3e-18
AH011815.2 Gorilla gorilla tyrosine hydroxylase (TH) gene, pa... https://www.ncbi.nlm.nih.gov/nuccore/AH011815.2	100	3e-18
XM_016149762.1 PREDICTED: Rousettus aegyptiacus insulin (INS)... https://www.ncbi.nlm.nih.gov/nuccore/XM_016149762.1	100	3e-18
XM_016149753.1 PREDICTED: Rousettus aegyptiacus insulin (INS)... https://www.ncbi.nlm.nih.gov/nuccore/XM_016149753.1	100	3e-18
KU548279.1 Uncultured bacterium clone PI_15F_Contig_5 genomic... https://www.ncbi.nlm.nih.gov/nuccore/KU548279.1	100	3e-18

KU548266.1	Uncultured bacterium clone CZ_15F_Contig_7 genomic...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/KU548266.1		
KU548224.1	Uncultured bacterium clone CH_15F_Contig_16 genomi...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/KU548224.1		
KU548210.1	Uncultured bacterium clone AZ_15F_Contig_2 genomic...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/KU548210.1		
NM_008386.4	Mus musculus insulin I (Ins1), mRNA	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/NM_008386.4		
XM_012743999.1	PREDICTED: Microcebus murinus insulin (INS), mRNA	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/XM_012743999.1		
XM_012651969.1	PREDICTED: Propithecus coquereli insulin (INS)...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/XM_012651969.1		
XM_007465953.1	PREDICTED: Lipotes vexillifer insulin (INS), mRNA	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/XM_007465953.1		
XM_006910977.1	PREDICTED: Pteropus alecto insulin (LOC1028811...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/XM_006910977.1		
XM_006750095.1	PREDICTED: Leptonychotes weddellii insulin (IN...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/XM_006750095.1		
XM_004317860.1	PREDICTED: Tursiops truncatus insulin (INS), mRNA	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/XM_004317860.1		
AB649280.1	Suncus murinus mRNA for insulin, partial cds	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/AB649280.1		
XM_002920120.1	PREDICTED: Ailuropoda melanoleuca insulin (INS...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/XM_002920120.1		
BC145868.1	Mus musculus insulin I, mRNA (cDNA clone MGC:17575...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/BC145868.1		
DQ250570.1	Niviventer coxingi preproinsulin 2 (Ins2) gene, co...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/DQ250570.1		
DQ250566.1	Niviventer coxingi preproinsulin 1 (Ins1) gene, co...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/DQ250566.1		
DQ250564.1	Apodemus semotus preproinsulin 1 (Ins1) gene, comp...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/DQ250564.1		
DQ250563.1	Rattus losea preproinsulin 1 (Ins1) gene, complete...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/DQ250563.1		
DQ479923.1	Mus musculus strain BTBR T+ tf/J insulin 1 precurs...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/DQ479923.1		
AC163452.12	Mus musculus chromosome 19, clone RP23-405C7, com...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/AC163452.12		
AK148541.1	Mus musculus adult pancreas islet cells cDNA, RIKE...	100	3e-18
	https://www.ncbi.nlm.nih.gov/nuccore/AK148541.1		

1.3 RISULTATI GRAFICI DELL'ANALISI

Attraverso il software della **T.T.E.S.**, la **sequenza acquisita** (o **sequenza originaria**) è sottoposta a un'analisi basata su uno dei suoi "Trend non manifesti": il **Trend n°1**.

Il grafico in Fig. 4 è relativo al **Profilo degli 8 Codici Principali** che descrivono *sinteticamente* la sequenza analizzata.

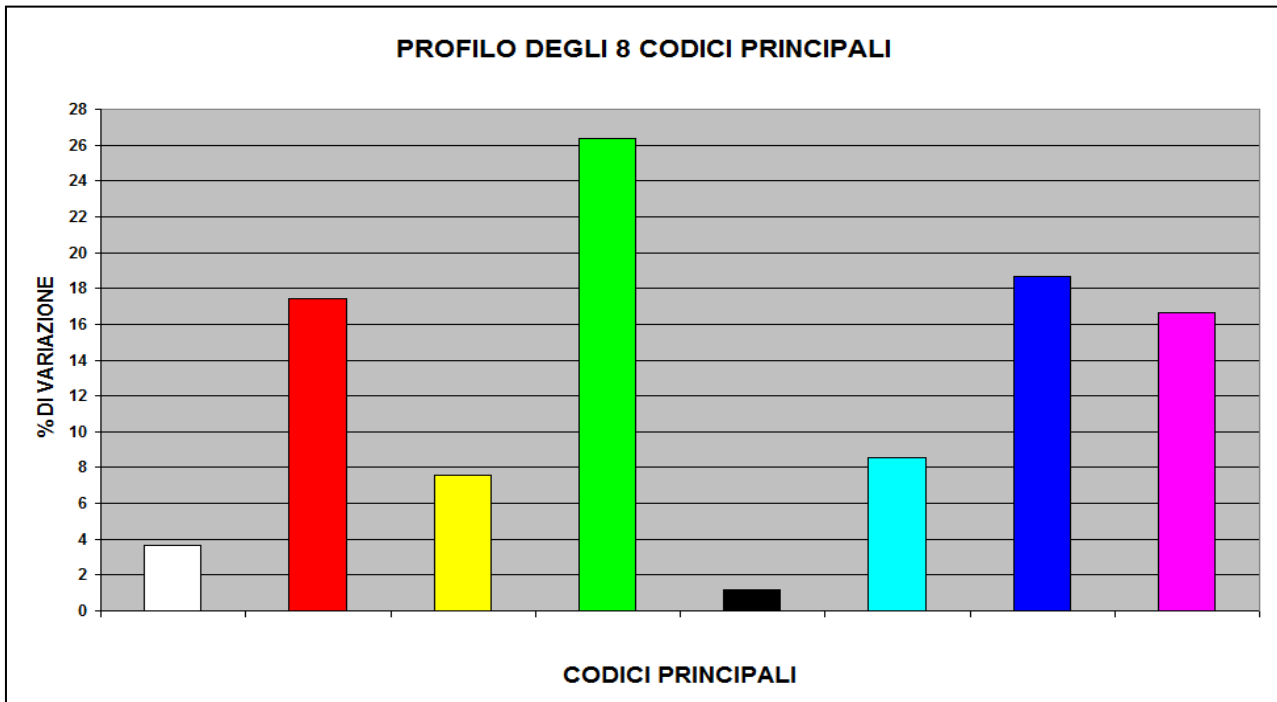


Fig. 4

Il grafico in Fig. 5 è relativo alla **Distribuzione della Percentuale di Variazione degli 8 Codici Principali** in riferimento alla sequenza analizzata.

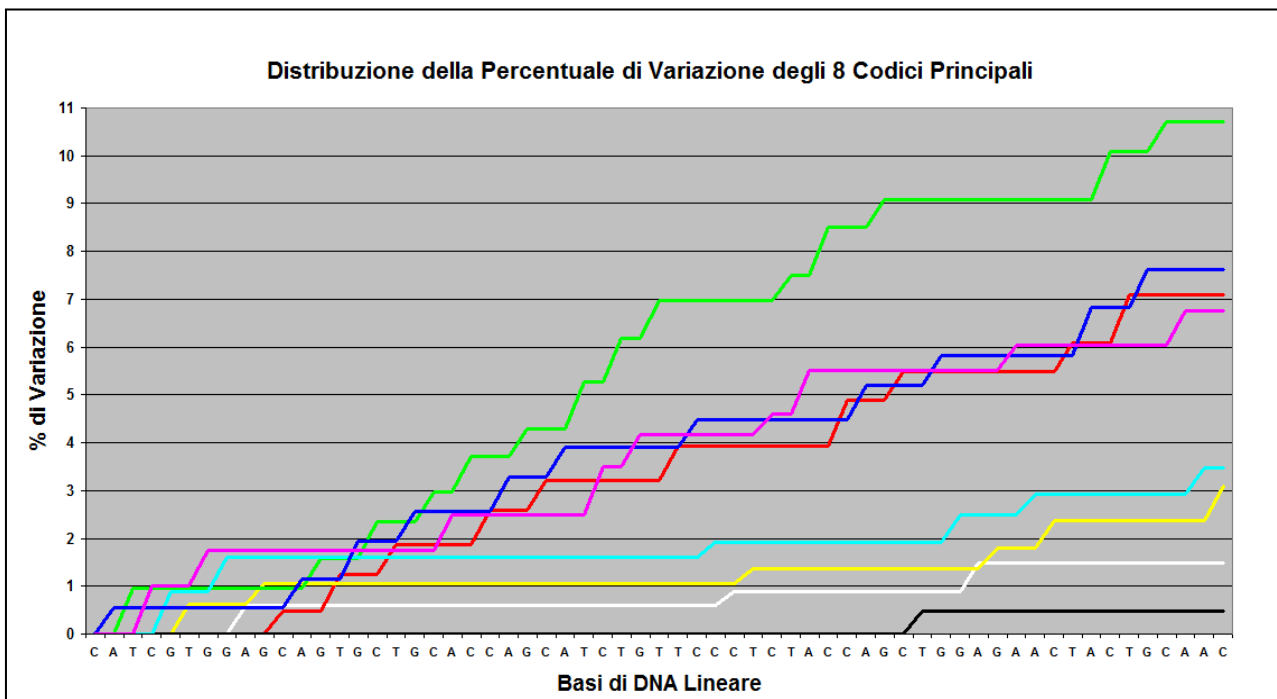


Fig. 5

Il grafico in Fig. 6 è relativo al profilo delle singole **Tonalità dei 64 Codici Totali** che descrivono *analiticamente* la sequenza analizzata.

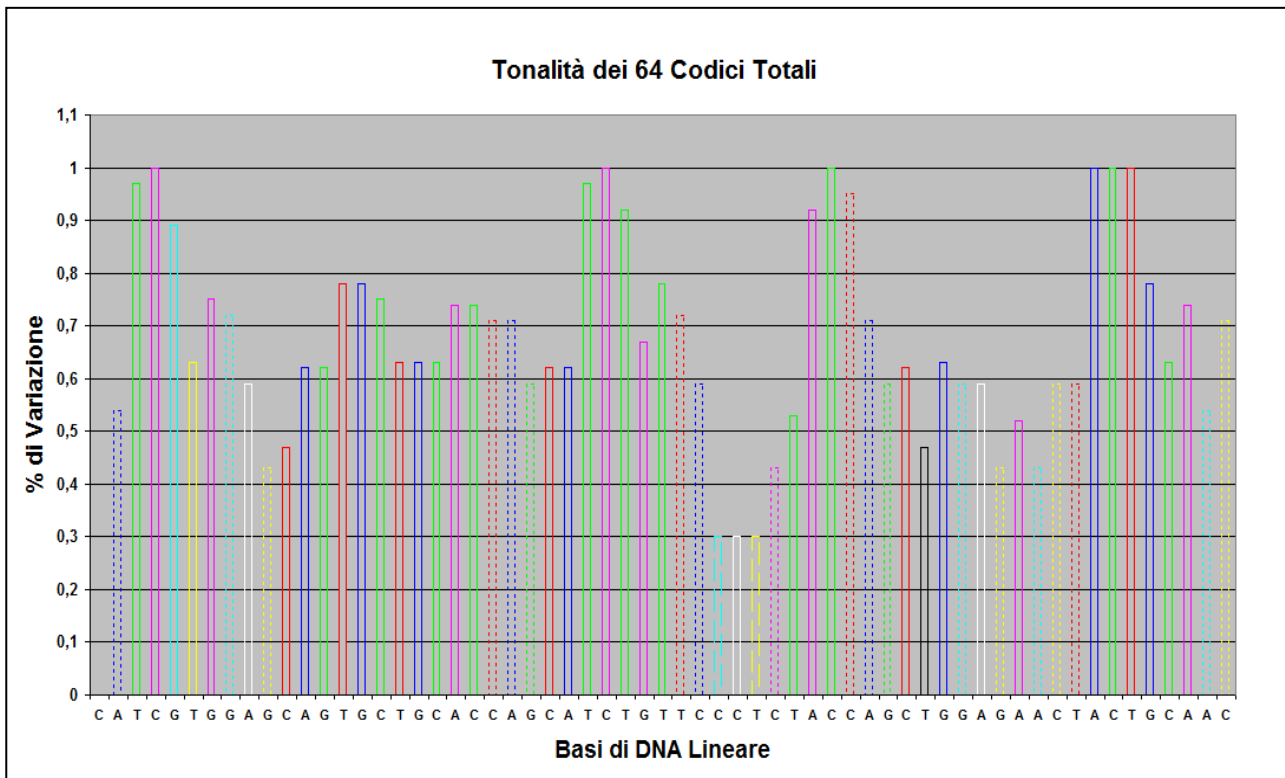


Fig. 6

Il grafico in Fig. 7 è relativo al **Profilo dei 64 Codici Totali** che descrivono *analiticamente* la sequenza analizzata.

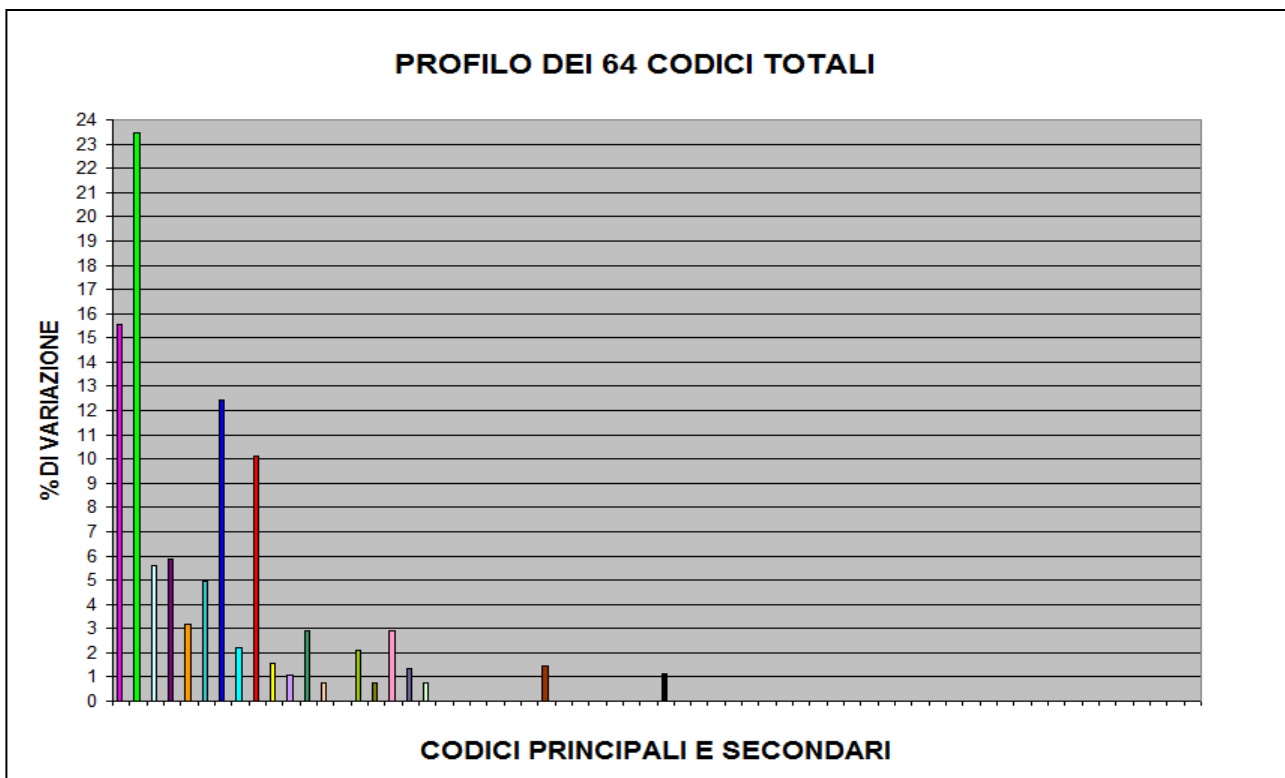


Fig. 7

1.4 RISULTATI GRAFICI DELL'ANALISI DELLA LETTURA NON CODIFICANTE E DI QUELLA CODIFICANTE DELLE BASI DI DNA

In Fig. 8 (A e B) sono confrontati due **Profili degli 8 Codici Principali**.

Nel grafico della Fig. 8 (A) la sequenza è analizzata **senza tenere conto** delle variazioni determinate dalla lettura CODIFICANTE (CDS <1.63) delle basi che codificano la **Catena A dell'Insulina**.

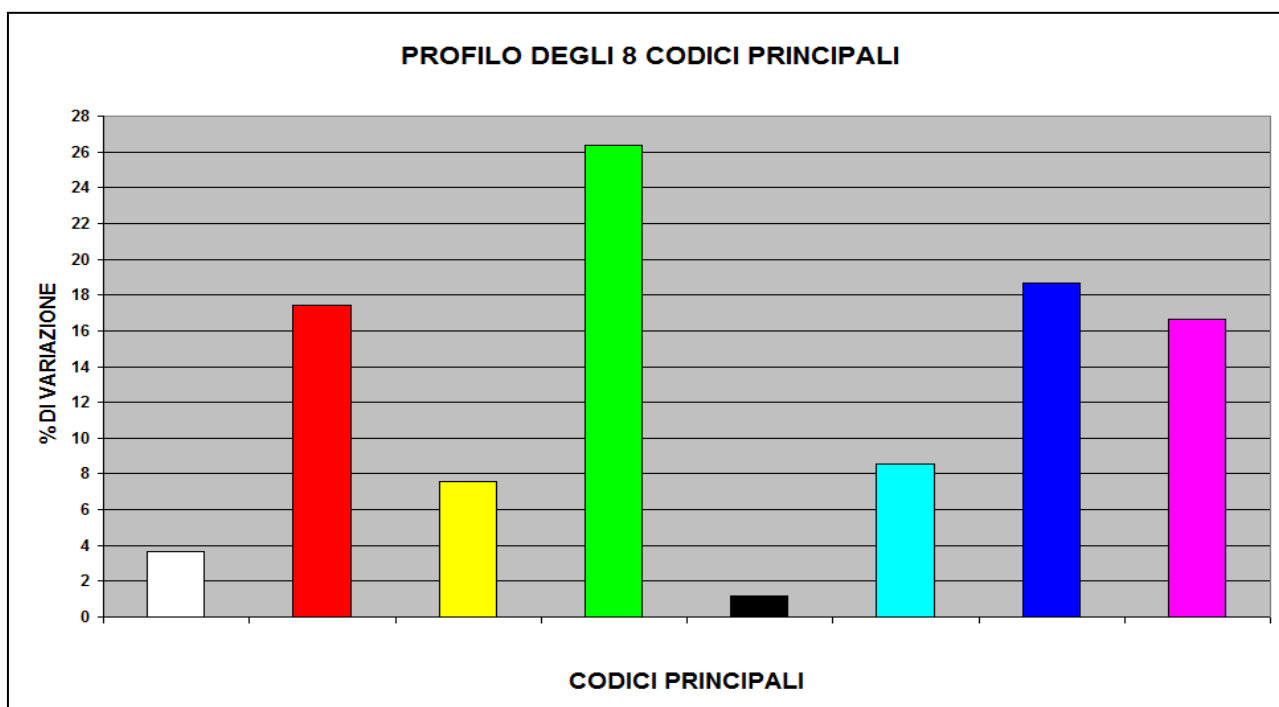


Fig. 8 (A)

Nel grafico in Fig. 8 (B) se ne tiene invece conto.

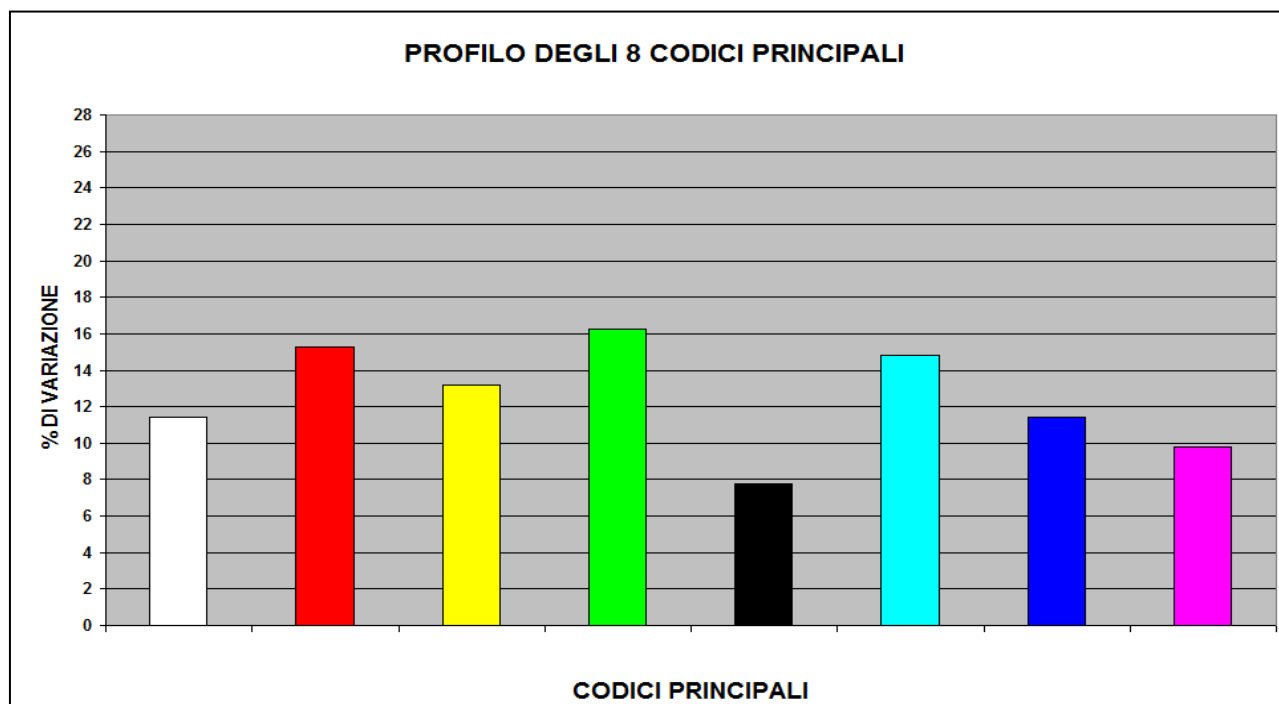


Fig. 8 (B)

In Fig. 9 (A e B) sono confrontati due grafici relativi alla **Distribuzione della Percentuale di Variazione degli 8 Codici Principali**.

Come nella figura precedente, nel grafico in Fig. 9 (A) **non si tiene conto** delle basi che CODIFICANO (CDS <1..63) la **Catena A dell'Insulina**.

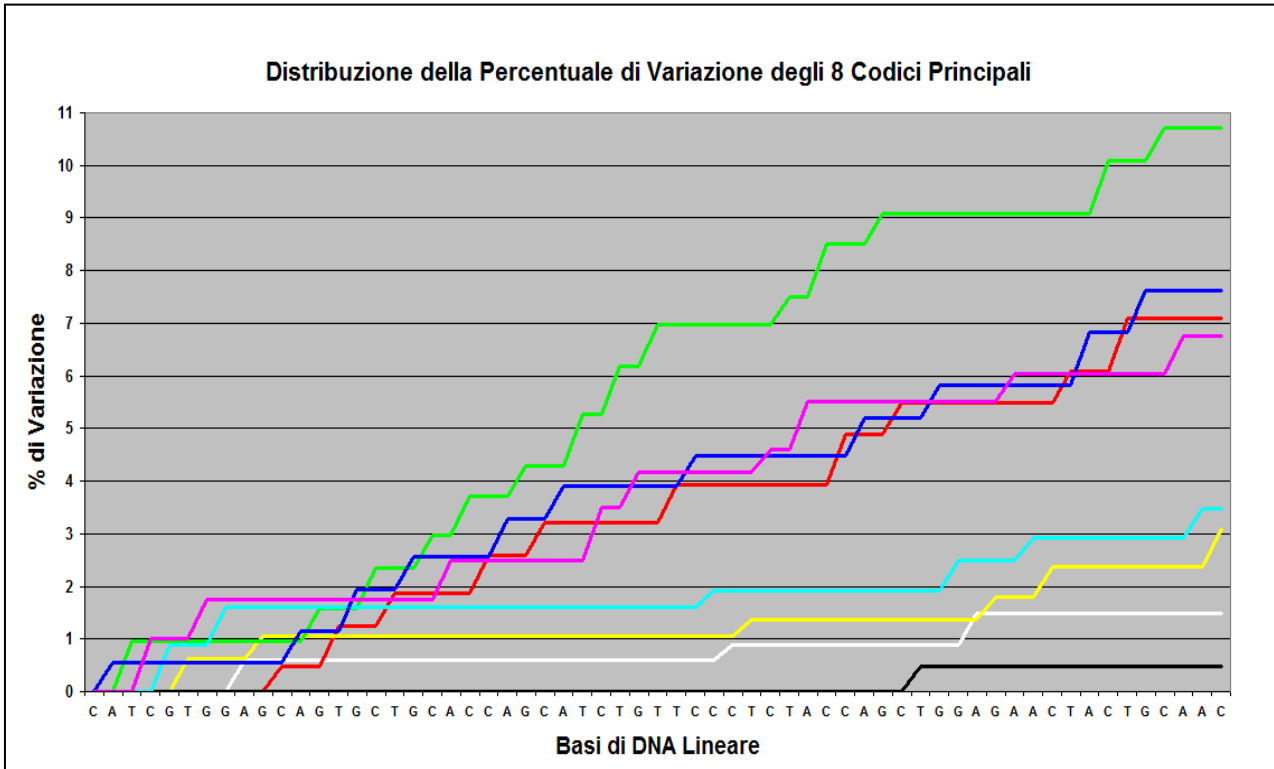


Fig. 9 (A)

Nel grafico in Fig. 9 (B) **se ne tiene invece conto**.

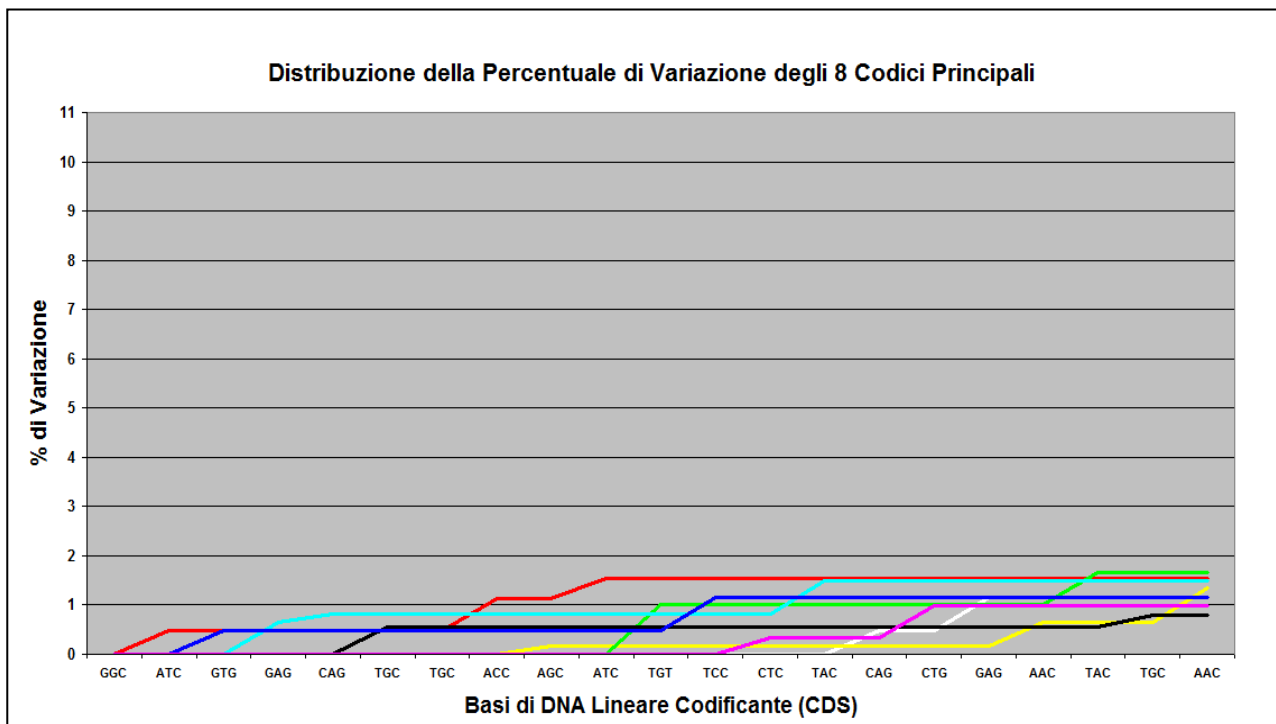


Fig. 9 (B)

In Fig. 10 (A e B) sono confrontati due grafici relativi alle singole **Tonalità dei 64 Codici Totali**.

Come nella figura precedente, nel grafico in Fig. 10 (A) **non si tiene conto** delle basi che CODIFICANO (CDS <1..63) la **Catena A dell'Insulina**.

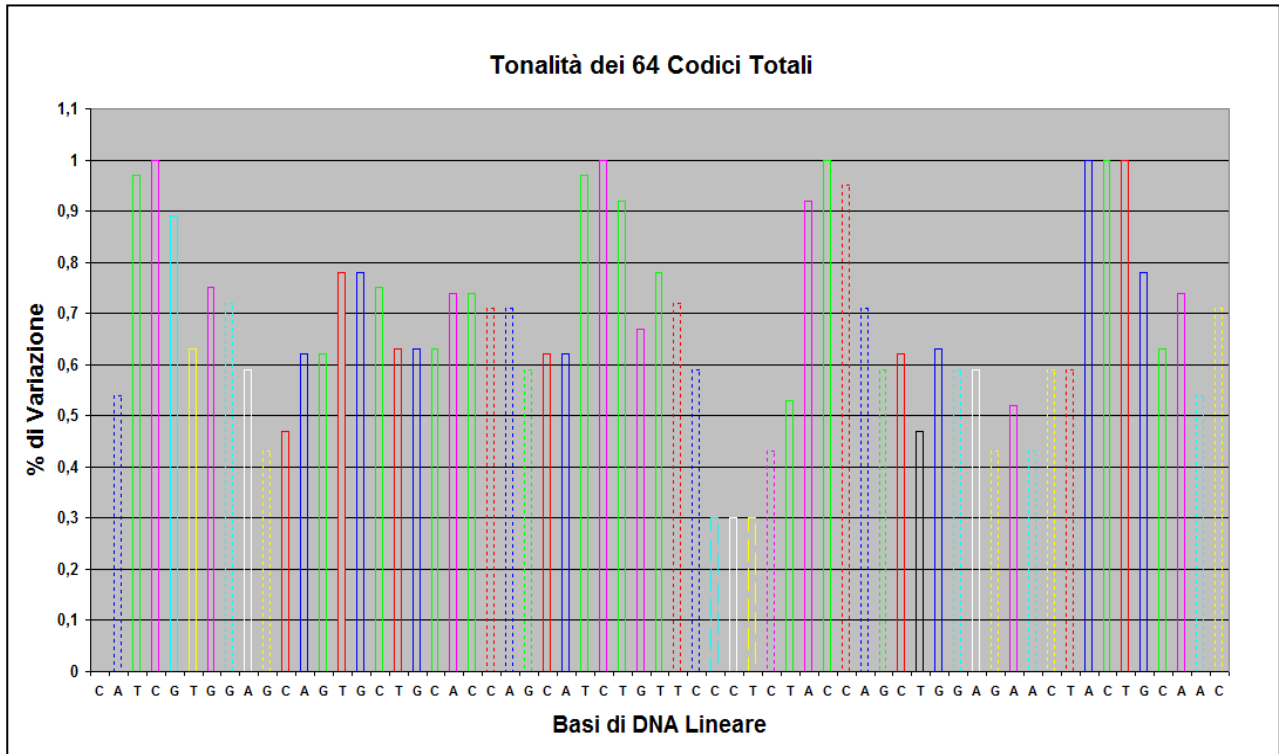


Fig. 10 (A)

Nel grafico in Fig. 10 (B) se ne tiene invece conto.

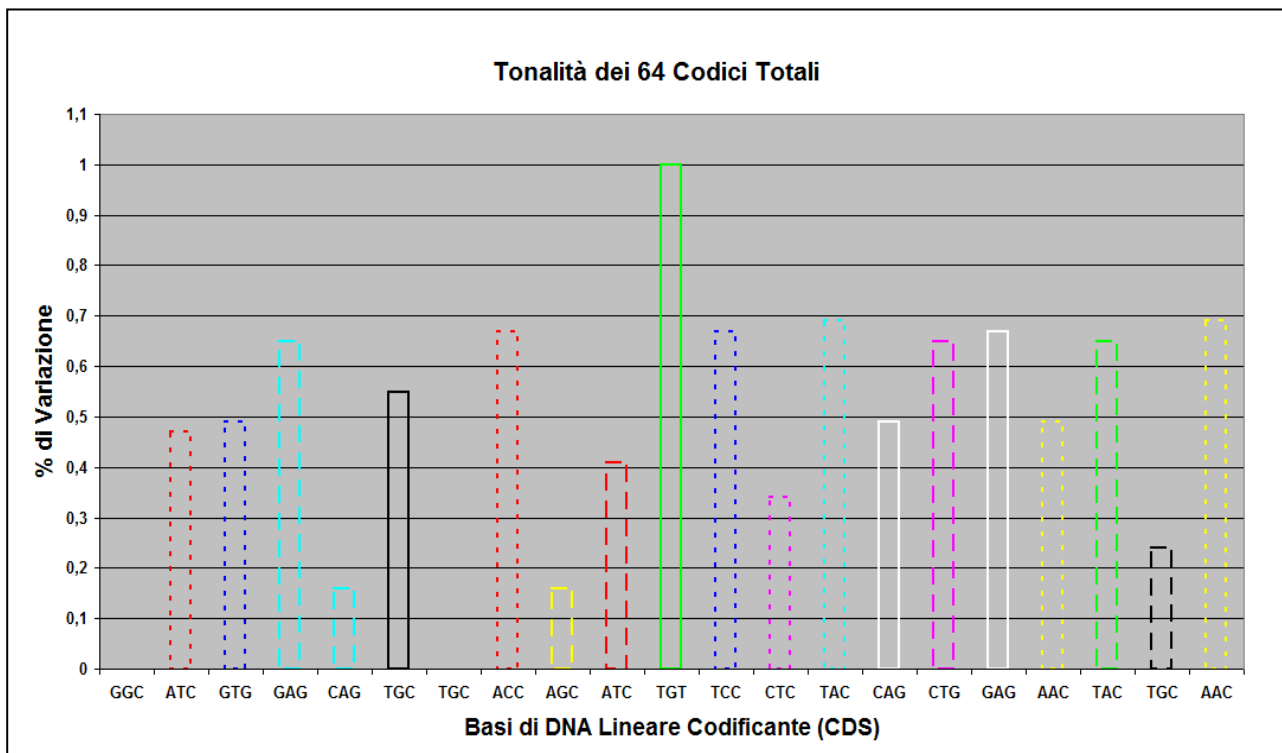


Fig. 10 (B)

In Fig. 11 (A e B) sono confrontati due grafici relativi al **Profilo dei 64 Codici Totali**.

Come nella figura precedente, nel grafico in Fig. 11 (A) **non si tiene conto** delle basi che CODIFICANO (CDS <1..63) la **Catena A dell'Insulina**.

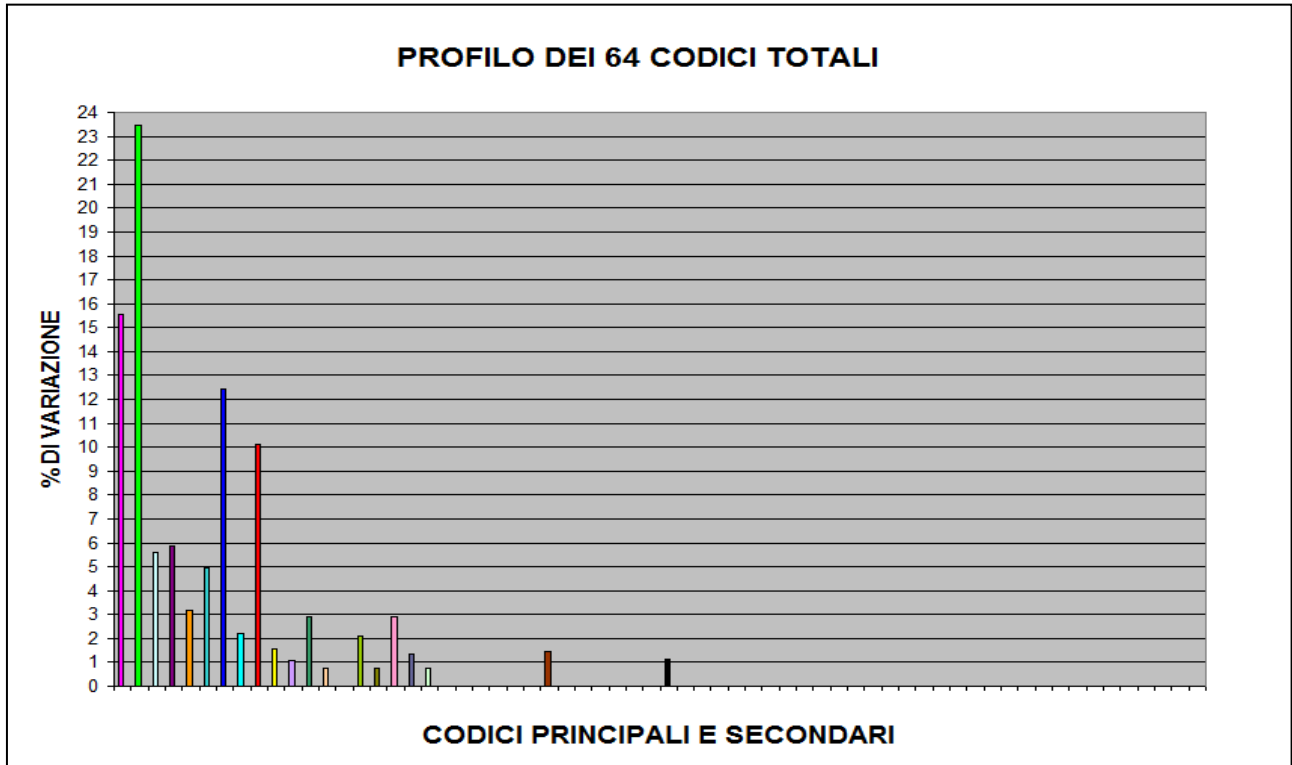


Fig. 11 (A)

Nel grafico in Fig. 11 (B) **se ne tiene invece conto**.

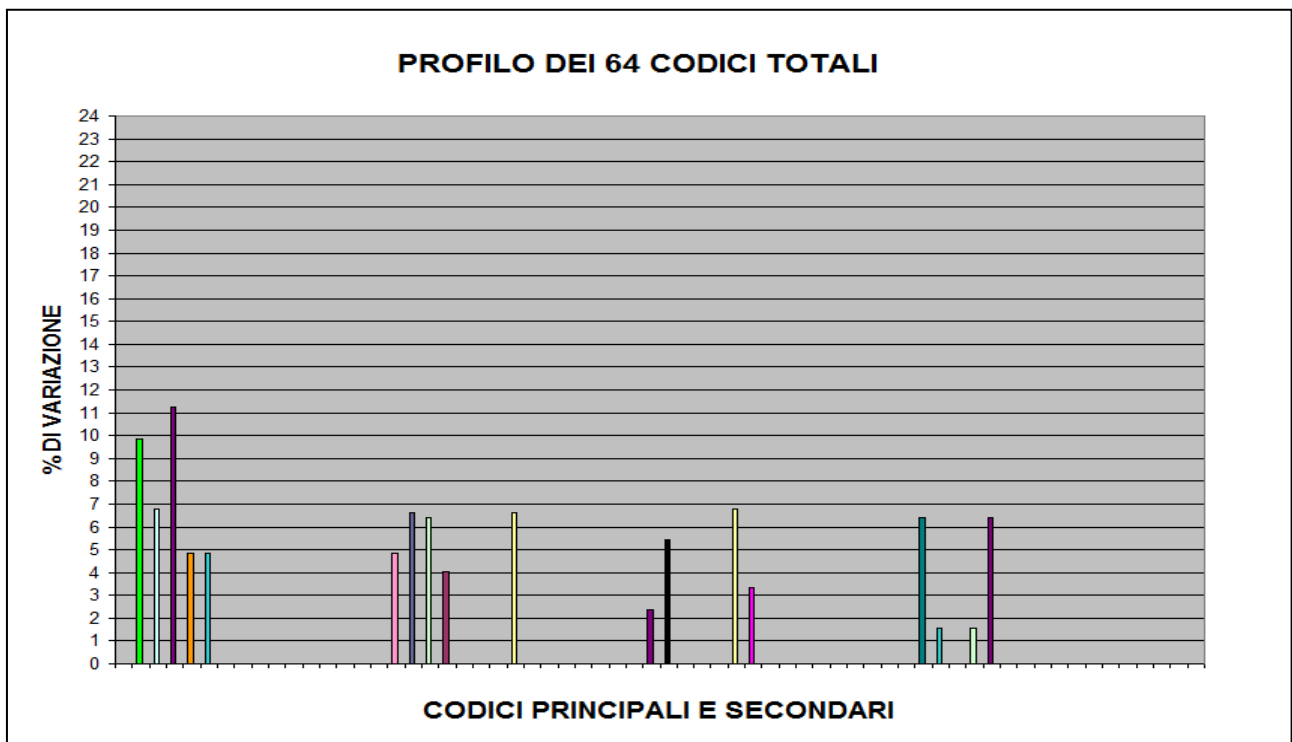


Fig. 11 (B)

**Analisi della
Sequenza n°1/1
della
Catena A dell'Insulina**

1.5 PREMESSE

Riconsideriamo la sequenza di **63** basi della **Catena A dell'Insulina** già analizzata:

ggcatcgtggagcagtgctgcaccagcatctgttcctctaccagctggagaactactgcaac

A partire da questa **sequenza originaria**, il software della T.T.E.S. genera numerose e diverse **nuove sequenze** che rispettano fedelmente i diversi e numerosi “**trend non manifesti**” della **sequenza originaria**.

Come è stato già accennato nell'**Introduzione**, in questo Capitolo, *che costituisce solo uno di molti altri che ne seguiranno*, delle numerose e diverse **nuove sequenze generate**, si è deciso di analizzarne solo una tra le possibili: la **Sequenza n°1/1**.

ATTENZIONE:

Per mantenere un interesse costante alla lettura complessiva di questo lavoro, nei risultati grafici qui di seguito presentati, sono state omesse alcune informazioni relative alla **nuova sequenza generata** (la **Sequenza n°1/1**).

Le informazioni totali riguardo tutte le **nuove sequenze generate** (compresa quindi la **Sequenza n°1/1**) e le loro corrispondenti *informazioni complete sugli allineamenti significativi* prodotti dalla *ricerca BLAST e i dati acquisiti da GenBank* (3) saranno pubblicate in **Appendice**, ovvero dopo la pubblicazione delle *Conclusioni generali*.

(3) Clark K., Karsch-Mizrachi I., Lipman D. J., Ostell J. and Sayers EW. GenBank. Nucleic Acids Res. 44(D1):D67-72 (2016). PMID: 26590407. PMCID: PMC4702903. <https://doi.org/10.1093/nar/gkv1276>

1.6 RISULTATI GRAFICI DELL'ANALISI DEL "TREND NON MANIFESTO"

Nella Fig. 12 (A e B) sono confrontati due **Profili degli 8 Codici Principali**.

Il grafico in Fig. 12 (A) si riferisce alla **sequenza originaria** di basi precedentemente analizzata.

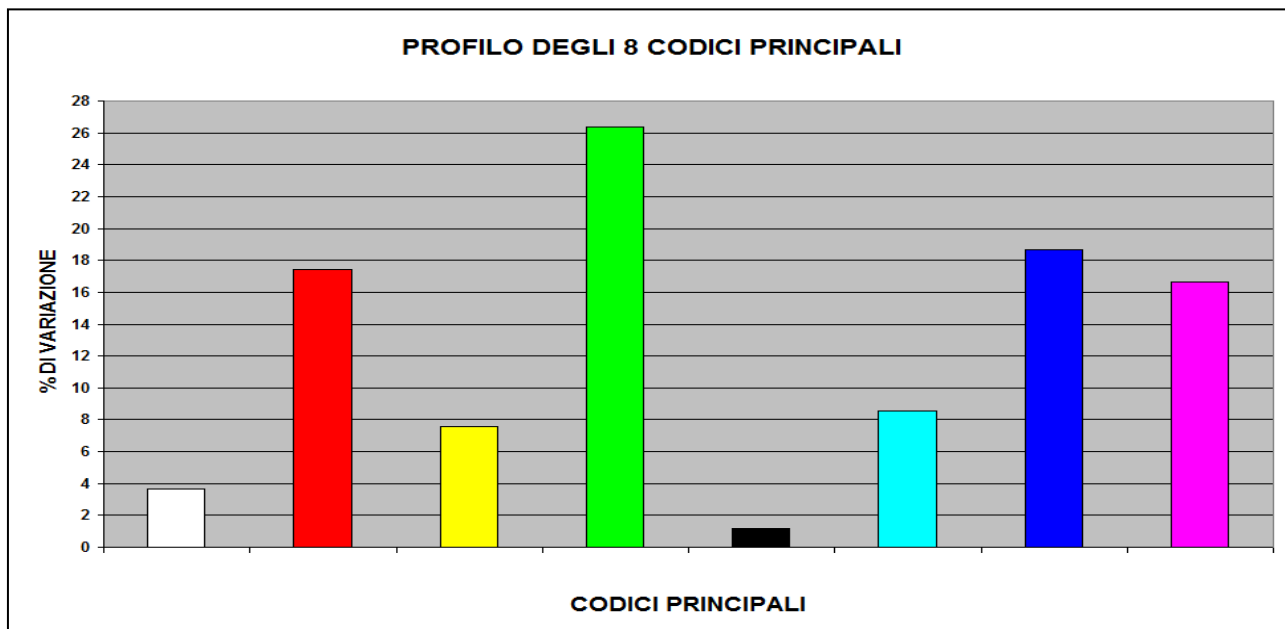


Fig. 12 (A)

Il grafico in Fig. 12 (B) si riferisce alla **"nuova sequenza generata"** a partire da quella originaria.

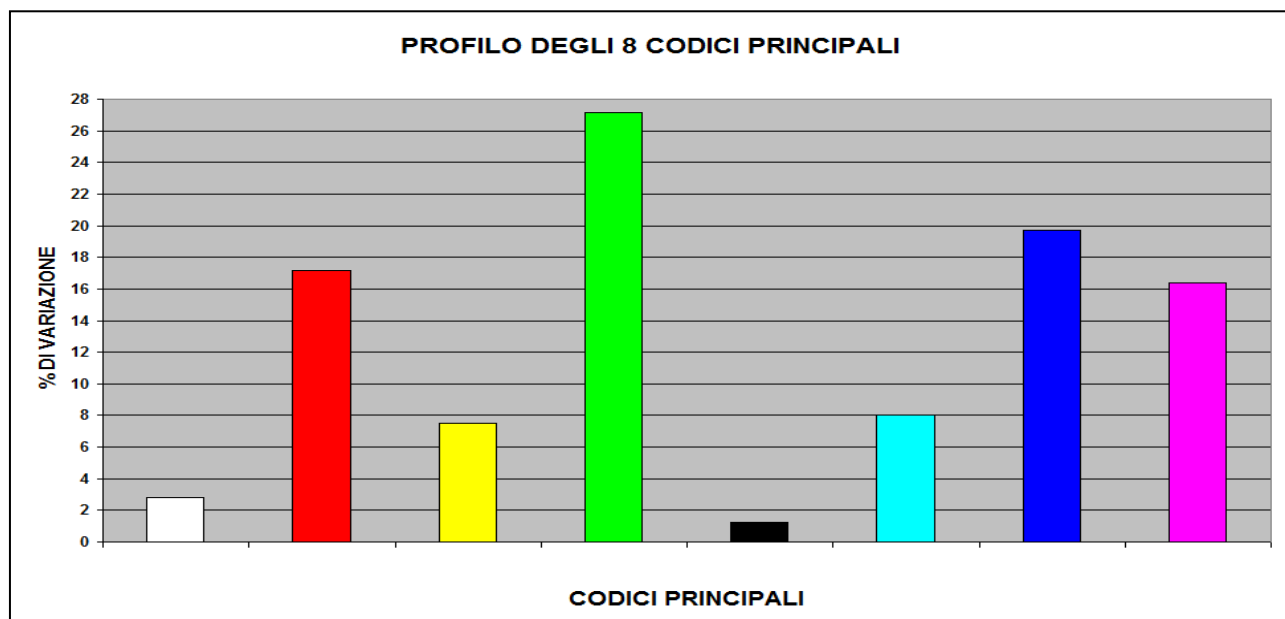


Fig. 12 (B)

Come si nota osservando i due grafici, i due **Profili degli 8 Codici Principali**, rispettivamente della **sequenza originaria** e della **nuova sequenza generata** a partire da quella originaria, SONO QUASI IDENTICI.

In Fig. 13 (A e B) sono confrontati due grafici relativi alla **Distribuzione della Percentuale di Variazione degli 8 Codici Principali**.

Il grafico in Fig. 13 (A) si riferisce alla **sequenza originaria** di basi precedentemente analizzata.

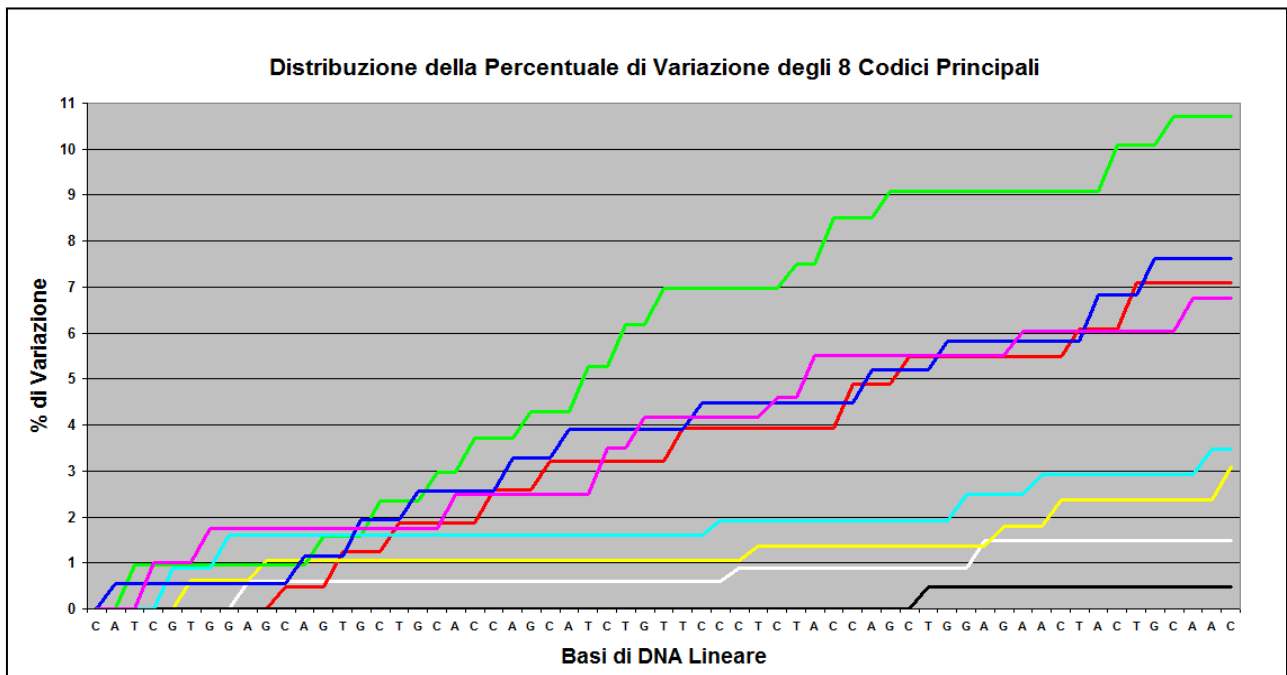


Fig. 13 (A)

Il grafico in Fig. 13 (B) si riferisce alla **“nuova sequenza generata”** a partire da quella originaria.

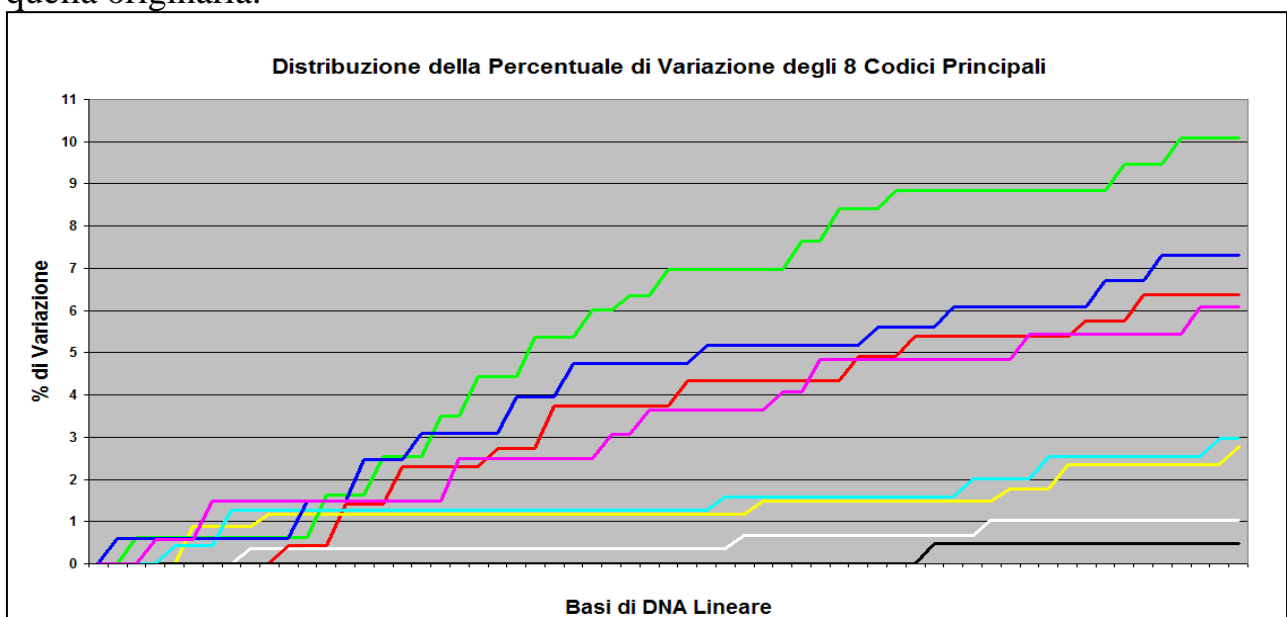


Fig. 13 (B)

Come per i due grafici della Fig. 12, anche in questo caso, i due grafici della **Distribuzione della Percentuale di Variazione degli 8 Codici Principali**, rispettivamente della **sequenza originaria** e della **nuova sequenza generata** a partire da quella originaria, SONO QUASI IDENTICI.

In Fig. 14 (A e B) sono confrontati due grafici relativi alle singole **Tonalità dei 64 Codici Totali**.

Il grafico in Fig. 14 (A) si riferisce alla **sequenza originaria** di basi precedentemente analizzata.

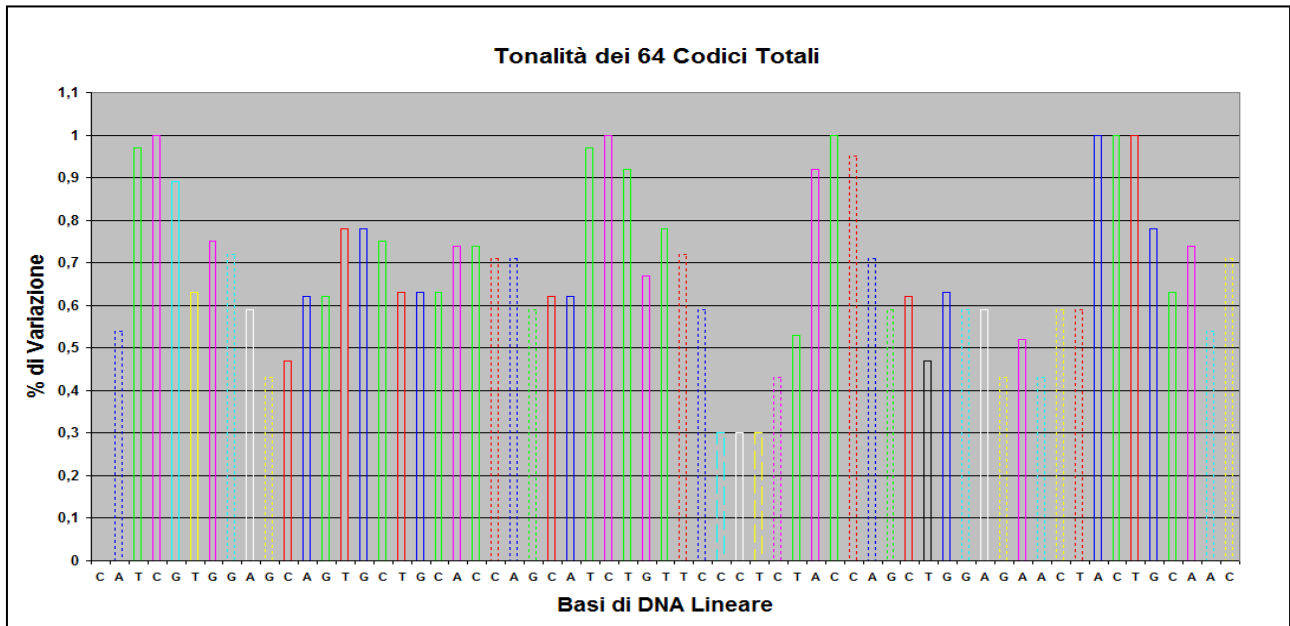


Fig. 14 (A)

Il grafico in Fig. 14 (B) si riferisce alla **“nuova sequenza generata”** a partire da quella originaria.

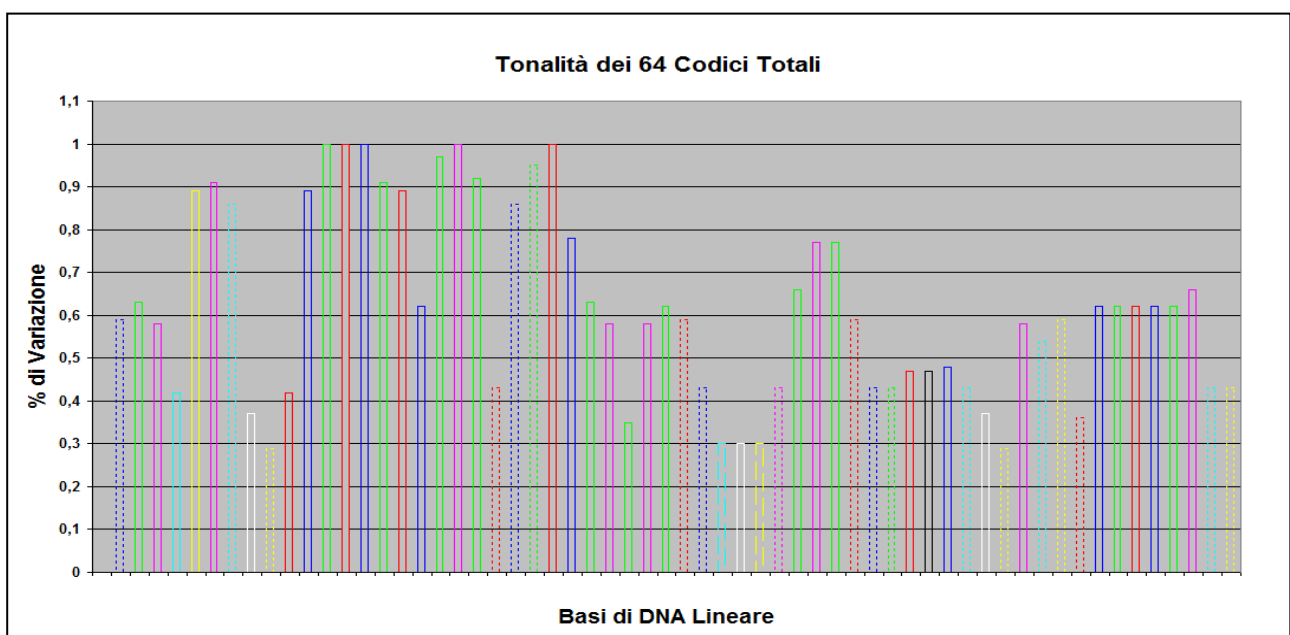


Fig. 14 (B)

A DIFFERENZA DEI GRAFICI PRECEDENTI, i due grafici delle singole **Tonalità dei 64 Codici Totali**, rispettivamente della **sequenza originaria** e della **nuova sequenza generata** a partire da quella originaria, SONO MOLTO DIVERSI TRA LORO.

In Fig. 15 (A e B) sono confrontati due grafici relativi al **Profilo dei 64 Codici Totali**.

Il grafico in Fig. 15 (A) si riferisce alla **sequenza originaria** di basi precedentemente analizzata.

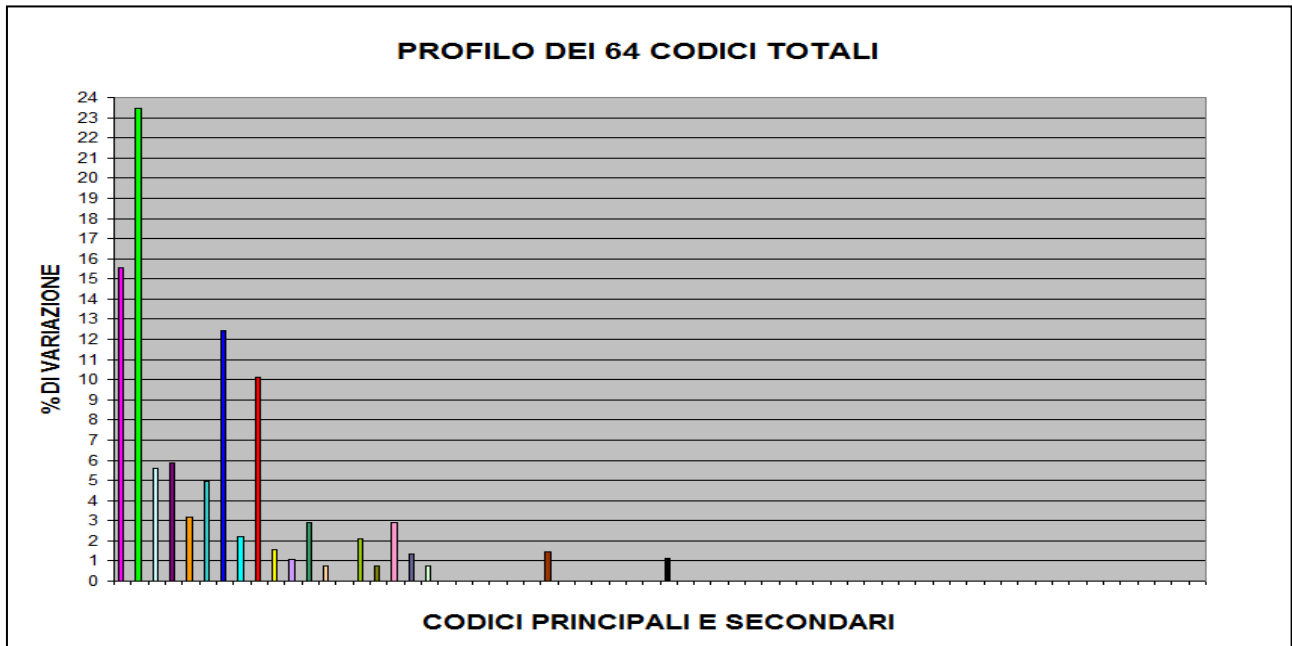


Fig. 15 (A)

Il grafico in Fig. 15 (B) si riferisce alla **“nuova sequenza generata”** a partire da quella originaria.

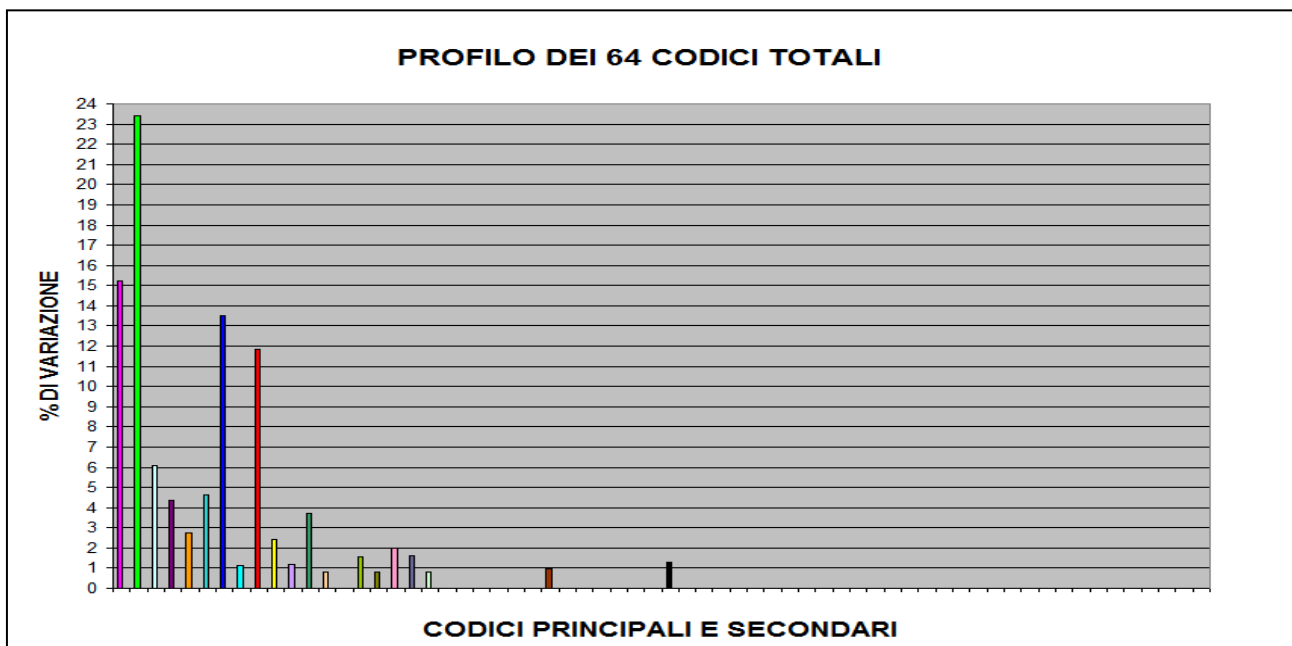


Fig. 15 (B)

A differenza dei due grafici precedenti, i due grafici relativi al **Profilo dei 64 Codici Totali**, rispettivamente della **sequenza originaria** e della **nuova sequenza generata** a partire da quella originaria, SONO IN PARTE SIMILI.

Dall'analisi della **T.T.E.S.** emerge che le differenze tra le “caratteristiche” della **sequenza originaria** e quelle della **nuova sequenza generata** a partire da quella originaria sono evidenziate in maniera rilevante dai grafici delle singole **Tonalità dei 64 Codici Totali** (Fig. 14 A e B).

Questo risultato è comprensibile se si considera che i grafici delle singole **Tonalità dei 64 Codici Totali** (Fig. 14 A e B), rispetto agli altri grafici, sono *molto più sensibili alle singole basi delle sequenze che al “trend non manifesto” delle sequenze stesse*.

Anche i grafici relativi ai **Profili dei 64 Codici Totali** (Fig. 15 A e B) evidenziano delle differenze, ma esse sono meno evidenti rispetto a quelle emerse dall'analisi dei grafici delle singole **Tonalità dei 64 Codici Totali** (Fig. 14 A e B).

Dagli altri grafici (Fig. 12 A e B, Fig. 13 A e B) emerge invece una quasi totale identità tra le “caratteristiche” della **sequenza originaria** e quelle della **nuova sequenza generata** a partire da quella originaria.

Questo risultato è notevole se si considera che **le 63 basi di DNA della nuova sequenza generata** sono QUASI TOTALMENTE DIVERSE DA QUELLE ORIGINARIE.

In conclusione, il grado di somiglianza delle “caratteristiche” della **nuova sequenza generata** alle “caratteristiche” della **sequenza originaria** è tanto maggiore quanto più simili sono rispettivamente i corrispondenti **Profili degli 8 Codici Principali** (Fig. 12 A e B) e quelli relativi alla **Distribuzione della Percentuale di Variazione degli 8 Codici Principali** (Fig. 13 A e B), *grafici entrambi molto sensibili al “trend non manifesto” delle sequenze*.

1.7 IMPLICAZIONI RELATIVE AI RISULTATI GRAFICI DELL'ANALISI DEL "TREND NON MANIFESTO"

Le **63** basi di DNA della **nuova sequenza generata** a partire da quella **originaria** sono state oggetto di ricerca **BLAST**.

Attenzione:

I risultati della *ricerca BLAST* saranno presentati gradualmente nei diversi capitoli che seguiranno. Nella parte prima del presente capitolo (Capitolo I°), sono presentati parzialmente solo i risultati relativi agli allineamenti con alcune specie di batteri **Pseudomonas** e con il nematoda **Heligmosomoides polygyrus**.

Le *informazioni totali sugli allineamenti significativi* prodotti dalla *ricerca BLAST* riguardo la **nuova sequenza generata (Sequenza n°1/1)** e le altre informazioni rilevanti ricavate da *GenBank* saranno pubblicate in **Appendice**, ovvero dopo la pubblicazione delle *Conclusioni generali*.

Query = NUOVA SEQUENZA GENERATA (Sequenza n°1/1)

Length = 63

Parametri della ricerca BLAST

Programma	<i>Blastn</i>
Word size	<i>11</i>
Expect value	<i>10</i>
Hitlist size	<i>100</i>
Match/Mismatch scores	<i>2,-3</i>
Gapcosts	<i>5,2</i>
Low Complexity Filter	<i>Yes</i>
Filter string	<i>L;m;</i>
Genetic Code	<i>1</i>

Sequences producing significant alignments:		Score	E	Identit.
		(Bits)	Value	
CP010359.1	<i>Pseudomonas plecoglossicida</i> strain NyZ12, complete...	44.6	0.18	85%
	https://www.ncbi.nlm.nih.gov/nuccore/CP010359.1			
LL188962.1	<i>Heligmosomoides polygyrus</i> genome assembly H_bak..	44.6	0.18	93%
	https://www.ncbi.nlm.nih.gov/nuccore/LL188962.1			
CP007620.1	<i>Pseudomonas putida</i> strain DLL-E4, complete genome	44.6	0.18	85%
	https://www.ncbi.nlm.nih.gov/nuccore/CP007620.1			
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LT629788.1	<i>Pseudomonas moraviensis</i> strain BS3668 genome ass....	39.2	7.6	82%
	https://www.ncbi.nlm.nih.gov/nuccore/LT629788.1			
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LL194531.1	<i>Heligmosomoides polygyrus</i> genome assembly H_bak..	39.2	7.6	87%
	https://www.ncbi.nlm.nih.gov/nuccore/LL194531.1			
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CP003961.1	<i>Pseudomonas</i> sp. VLB120, complete genome	39.2	7.6	83%
	https://www.ncbi.nlm.nih.gov/nuccore/CP003961.1			
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ALLINEAMENTI SIGNIFICATIVI

1.8 ALLINEAMENTI PSEUDOMONAS

>CP010359.1

Pseudomonas plecoglossicida strain NyZ12, complete genome

Length=6233254

Features in this part of subject sequence: **cystathionine gamma-synthase**

<https://www.ncbi.nlm.nih.gov/nucleotide/752308899?report=gbwithparts&from=5560737&to=5562680&RID=1ZXSZEJC014>

>CP007620.1

Pseudomonas putida strain DLL-E4, complete genome

Length=6484062

Features in this part of subject sequence: **cystathionine gamma-synthase**

<https://www.ncbi.nlm.nih.gov/nucleotide/635291785?report=gbwithparts&from=176815&to=178758&RID=1ZXSZEJC014>

>LT629788.1

Pseudomonas moraviensis strain BS3668 genome assembly, chromosome:I

Length=6092541

Features in this part of subject sequence: **high-affinity iron transporter**

<https://www.ncbi.nlm.nih.gov/nucleotide/1086004611?report=gbwithparts&from=4649680&to=4651578&RID=1ZXSZEJC014>

>CP003961.1

Pseudomonas sp. VLB120, complete genome

Length=5644569

Features in this part of subject sequence: **cytochrome c class I**

<https://www.ncbi.nlm.nih.gov/nucleotide/556072477?report=gbwithparts&from=5441006&to=5442901&RID=1ZXSZEJC014>

1.9 ALLINEAMENTI HELIGMOSOMOIDES POLYGYRUS

>LL188962.1

Heligmosomoides polygyrus genome assembly H_bakeri_Edinburgh, scaffold HPBE_scaffold0000593

Length=94530

[https://www.ncbi.nlm.nih.gov/nucleotide/688429340?report=genbank&log\\$=nuclalig n&blast_rank=2&RID=27MWTXV3014](https://www.ncbi.nlm.nih.gov/nucleotide/688429340?report=genbank&log$=nuclalig n&blast_rank=2&RID=27MWTXV3014)

>LL194531.1

Heligmosomoides polygyrus genome assembly H_bakeri_Edinburgh, scaffold HPBE_contig0000102

Length=27221

[https://www.ncbi.nlm.nih.gov/nucleotide/688443549?report=genbank&log\\$=nuclalig n&blast_rank=23&RID=27MWTXV3014](https://www.ncbi.nlm.nih.gov/nucleotide/688443549?report=genbank&log$=nuclalig n&blast_rank=23&RID=27MWTXV3014)

1.10 PSEUDOMONAS

La ricerca BLAST evidenzia *allineamenti significativi di basi di DNA* tra la **nuova sequenza generata** e diverse specie del genere dei **batteri PSEUDOMONAS**.

[**Pseudomonas** è un genere di batteri (Gram-negativi, aerobi obbligati ossidasi positivi, catalasi positivi e capaci di muoversi) appartenenti alla famiglia delle *Pseudomonadaceae*. Sono presenti soprattutto nel terreno, negli ambienti umidi e nelle acque (piscine) e sulle piante. Producono la *piocianina* (un tipo di pigmento) che altera la funzione ciliare, stimola la risposta infiammatoria e provoca danno ai tessuti. Una particolare sostanza prodotta da molte specie di *Pseudomonas* in condizioni di carenza di ferro è la *pioverdina* (un *sideroforo giallo-verde* o molecola ad alta affinità per il ferro capace di chelarlo efficacemente). I principali fattori di virulenza sono l'*esotossina A* e l'*endotossina*. Aderendo alle cellule dell'ospite possono formare *biofilm* (biopellicole protette da matrici di esopolisaccaridi) che inibiscono la fagocitosi e limitano l'attività antibiotica. Nell'uomo, la specie ***Pseudomonas aeruginosa*** è probabilmente la più diffusa (feci, ma anche ascelle, inguine e, più raramente, unghie). Nei nosocomi con carenze igieniche possono aversi epidemie con conseguenze potenzialmente gravi (infezioni osteoarticolari, otite esterna, polmonite, follicoliti cutanee, infezioni oculari ed endocardite). *Essendo poco permeabili, resistono alla maggior parte degli antibiotici o li espellono, inattivano le penicilline e gli aminoglicosidi producendo enzimi specifici*. Recentemente, per contrastare la virulenza di *Pseudomonas*, i ricercatori hanno sfruttato la sua forte avidità per il ferro. Gli ioni di gallio interagiscono con i processi cellulari in modo simile al ferro (III). Quando batteri come *Pseudomonas* scambiano per errore gli ioni di gallio con il ferro (III), gli ioni interferiscono con la respirazione e i batteri muoiono. Ciò accade perché il **ferro è attivo redox**, consentendo il trasferimento di elettroni durante la respirazione, mentre il **gallio è inattivo redox**](4).

Il **primo allineamento significativo** della ricerca BLAST riguarda il **batterio PSEUDOMONAS** della specie *Plecoglossicida*.

Dalla stessa ricerca emergono successivi *altri allineamenti significativi* con altre 3 specie di **batteri PSEUDOMONAS**: *Putida*, *Sp.* e *Moraviensis*.

In particolare, nei **batteri PSEUDOMONAS** delle specie ***Plecoglossicida Nyz12*** e ***Putida DLL-E4*** gli *allineamenti di basi* hanno riguardato l'**enzima CYSTATHIONINE GAMMA-SYNTHASE**.

Invece, nei **batteri PSEUDOMONAS Moraviensis BS3668** l'*allineamento* ha evidenziato un HIGH-AFFINITY IRON TRANSPORTER, mentre nei **batteri PSEUDOMONAS** della specie ***Sp. VLB120*** l'*allineamento* di basi ha riguardato il CYTOCHROME C CLASS 1.

(4) Tratto e modificato da: <https://it.wikipedia.org/wiki/Pseudomonas>

<https://en.wikipedia.org/wiki/Pseudomonas>

1.11 HELIGMOSOMOIDES POLYGYRUS

[*Heligmosomoides polygyrus* (conosciuto anche come *Nematospiroides dubius*) è un nematoda intestinale dei roditori selvatici (in particolare, dei topi selvatici *Apodemus sylvaticus*). Esso ha la capacità di stabilire infezioni croniche nei roditori e alterarne le risposte immunitarie. Per questo motivo, *Heligmosomoides polygyrus* è ampiamente usato (soprattutto sul topo di laboratorio *Mus musculus*) come modello parassitario gastrointestinale in studi immunologici, farmacologici e tossicologici. Il roditore infetto scatena una forte risposta immunitaria innata e adattiva, generata per prevenire la formazione del parassita nell'intestino e per guarire le ferite (immunità tipo Th2) associate alla patologia intestinale. Tuttavia, nonostante questa impressionante risposta immunitaria, *H. polygyrus* è in grado di dirottare la risposta immunitaria dell'ospite, di attenuare la risposta Th2 e di causare un'infezione cronica] (5).

(5) Tratto e modificato da: https://en.wikipedia.org/wiki/Heligmosomoides_polygyrus

1.12 PRESUPPOSTI PER LE RICERCHE BIBLIOGRAFICHE MIRATE

In questo capitolo, come negli altri capitoli che seguiranno, saranno fatte delle **ricerche bibliografiche mirate** finalizzate a trovare le possibili relazioni tra l'**insulina** e i diversi e numerosi *organismi* identificati dalle ricerche *BLAST* sulle diverse **nuove sequenze generate**.

Nella prima parte di questo capitolo, le ricerche riguarderanno i batteri del genere **Pseudomonas** e il nematoda **Heligmosomoides polygyrus**.

Nella seconda parte di questo capitolo, le ricerche saranno effettuate su tutti i restanti *organismi* per i quali sono stati riscontrati allineamenti significativi con la **Sequenza n°1/1**.

I risultati delle ricerche bibliografiche saranno consultabili direttamente dai **link ad abstract di articoli** (o a **interi articoli**) **scientifici** pubblicati su **PubMed** (<https://www.ncbi.nlm.nih.gov/pubmed/>), [una risorsa gratuita sviluppata e gestita dal **NCBI** (National Center for Biotechnology Information), presso la **NLM** (National Library of Medicine) degli Stati Uniti, situata presso il **NIH** (National Institutes of Health)](si veda anche *PubMed Help: How to Get the Journal Article*).

La trattazione degli articoli esula dalle finalità del lavoro qui proposto e, dato che non sempre dal titolo dell'articolo si evincono le importanti relazioni, dirette o indirette, tra l'insulina e i diversi organismi, si rimanda il lettore interessato al loro studio e approfondimento.

Il principale obiettivo, nell'elencare questi articoli scientifici, è quello di accumulare prove (e materiale da approfondire) che sostengano come scientificamente plausibile l'*ipotesi* che le **nuove sequenze generate** abbiano ragionevoli **forti relazioni** con le **caratteristiche** della **sequenza originaria** (nello specifico con la **Catena A dell'Insulina**).

Questa ipotesi sarebbe sostenibile e verosimile se le ricerche bibliografiche confermassero che la **sequenza originaria** fosse in qualche modo implicata con alcune caratteristiche degli *organismi* costituiti da basi di DNA (o RNA) delle diverse nuove sequenze generate.

1.13 LA RICERCA RIGUARDANTE PSEUDOMONAS

I risultati della ricerca BLAST (presentati a pagina 33) hanno stimolato un'opportuna ricerca bibliografica per studiare le **possibili relazioni** tra le **caratteristiche** della **sequenza originaria di DNA** (**63** basi della **Catena A dell'Insulina**) e le **caratteristiche** dei batteri **Pseudomonas**.

Dalla lettura degli articoli selezionati (presentati dalla pagina 51 alla pagina 156) sono state riscontrate importanti relazioni, *dirette* o *indirette*, tra l'**insulina** e diverse specie di batteri del genere **Pseudomonas**.

Queste forti relazioni tra l'**insulina** e **Pseudomonas** avvalorano l'ipotesi che la **Sequenza n°1/1**, cioè la **nuova sequenza generata** rispettando fedelmente il **Trend n°1** (che è uno tra i possibili specifici "trend non manifesti" della *sequenza originaria*), abbia **forti relazioni** con le **caratteristiche** della **sequenza originaria** (nello specifico con la **Catena A dell'Insulina**).

In generale e molto sommariamente, tra le altre prove di minore importanza non citate, **esistono prove di forti legami** tra *Insulina*, *Pseudomonas* (o *Burkholderia*, una specie del genere di batteri *Pseudomonas*) e:

- 1) *Diabete*;
- 2) *Melioidosi*;
- 3) *Fibrosi Cistica*;
- 4) *Infezioni Polmonari*;
- 5) *Obesità*;
- 6) *Otite Esterna Maligna*;
- 7) *Endocardite*;
- 8) *Sistema Immunitario*;
- 9) *Apoptosi*.

Più precisamente, *Pseudomonas* è molto più presente e causa conseguenze molto più gravi [tra le quali l'*Otite Esterna Maligna* e l'*Endocardite* (quest'ultima soprattutto nei tossicodipendenti)] in soggetti immunodepressi e/o affetti da *Diabete*, *Fibrosi Cistica*, *Infezioni Polmonari* e *Obesità*, che in soggetti che non sono affetti da queste patologie. L'attività infettiva di *Pseudomonas* può indebolire le difese immunitarie dell'ospite e favorire il processo dell'*apoptosi* (morte cellulare programmata) e il cancro.

La **melioidosi** è invece un'infezione causata dal batterio *Burkholderia* (o *Pseudomonas*) *pseudomallei*. Anch'essa è molto più presente e grave quando i soggetti sono affetti soprattutto da *Diabete*, *Fibrosi Cistica* e *Infezioni Polmonari*.

1.14 CYSTATHIONINE GAMMA-SYNTHASE

Il risultato della ricerca BLAST identifica allineamenti significativi di basi con l'enzima CYSTATHIONINE GAMMA-SYNTHASE dei batteri **Pseudomonas**. Questo risultato ha stimolato una **ricerca bibliografica specifica** sulle relazioni tra **Insulina, Pseudomonas e Cystathionine Gamma-Synthase**.

[L'enzima **CYSTATHIONINE GAMMA-SYNTHASE (CGS)** è un enzima che catalizza una reazione chimica. I due substrati di questo enzima sono O 4 succinyl-L-homoserine + L-cysteine, mentre i suoi due prodotti sono L-cystathionine + succinate. Questo enzima partecipa a 4 percorsi metabolici: il *metabolismo della metionina*, il *metabolismo della cisteina*, il *metabolismo dei seleno-aminoacidi* e il *metabolismo dello zolfo*](6).

La ricerca bibliografica su tutti gli **articoli scientifici** pubblicati su **PubMed** in cui figurano insieme i termini “*Insulin*”, “*Pseudomonas*” e “*Cystathionine Gamma-Synthase*” **non ottiene risultati**.

(<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+PSEUDOMONAS+CYSTATHIONINE+GAMMA-SYNTHASE>).

Questa constatazione fa ipotizzare possibili **relazione indirette** tra *Insulin*, *Pseudomonas* e *Cystathionine Gamma-Synthase*.

Per indagare su queste **ipotetiche relazioni** si è proseguito facendo delle **ricerche mirate** sugli accoppiamenti dei termini “*Insulin e Cystathionine Gamma-Synthase*” e “*Pseudomonas e Cystathionine Gamma-Synthase*”.

Dagli articoli che riguardano la **ricerca mirata** sugli accoppiamenti dei termini “*Insulin e Cystathionine Gamma-Synthase*” emergono principalmente **prove di forti legami** tra *Insulin*, *Cystathionine Gamma-Synthase* e:

- 1) *Diabete*;
- 2) *Obesità*;
- 3) *Solfato di Idrogeno (H₂S)*;
- 4) *Omocisteina*;
- 5) *Cisteina*.

(6) Tratto e modificato da:

https://en.wikipedia.org/wiki/Cystathionine_gamma-synthase

Fondamentale appare il ruolo dell'**Omocisteina** e della **Cisteina** e dei loro rapporti con il **Solfato di Idrogeno** (sintetizzato nell'uomo soprattutto dagli enzimi *cistationina- γ -liasi* e *cistationina- β -sintasi* degli amminoacidi cistationina, omocisteina e cisteina) nel **Diabete** e nell'**Obesità**.

Esiste quindi un evidente legame (diretto e indiretto) tra l'**Insulina** e più di un **percorso metabolico in cui è implicato l'enzima CYSTATHIONINE GAMMA-SYNTASE** (metabolismo della *metionina*, metabolismo della *cisteina*, metabolismo dei *seleno-aminoacidi* e metabolismo dello *zolfo*).

In particolare, il coinvolgimento del **Solfato di Idrogeno** sposta l'attenzione sulla sua attività **antiossidante** e sulla sua capacità di influenzare lo **stato redox delle cellule** [inibendo l'azione fisiologicamente disfunzionale delle **ROS** (*Reactive Oxygen Species*), Specie chimiche **Reattive** chimicamente contenenti **Ossigeno**, quali i *peroxides*, *superoxide*, *hydroxyl radical* e *singlet oxygen*].

[Nei processi biologici, le **ROS** sono formate come sottoprodotto naturale del normale metabolismo dell'**ossigeno** e hanno ruoli importanti nella **segnalazione cellulare** e nell'**omeostasi**. In seguito allo **stress ossidativo** (dovuto a vari fattori ambientali), i *livelli delle ROS* possono aumentare molto e causare danni significativi alle strutture cellulari. Le ROS sono prodotti intracellulari e le fonti principali sono i complessi enzimatici **NADPH ossidasi** (NOX) presenti nelle *membrane cellulari*, nei *mitocondri*, nei *perossisomi* e nel *reticolo endoplasmatico*. Nei mitocondri, la **fosforilazione ossidativa**, comporta il trasporto di protoni (ioni idrogeno) mediante la **catena di trasporto degli elettroni**. L'ultima destinazione di un elettrone lungo questa catena è una molecola di *ossigeno*. In condizioni normali, l'ossigeno viene ridotto per produrre acqua; tuttavia, in una percentuale bassissima di elettroni che passano attraverso la catena, l'ossigeno è invece prematamente e in modo incompleto ridotto e si ottiene un radicale superossido. Esso può inattivare specifici enzimi o iniziare la perossidazione lipidica nella sua forma protonata, producendo dei danni. Se nei mitocondri è presente un danno eccessivo, la *cellula* subisce l'**apoptosi** (*morte cellulare programmata*). Le **proteine Bcl-2** sono stratificate sulla superficie dei mitocondri, rilevano il danno e attivano una classe di proteine chiamate **Bax** che causando la fuoriuscita del **Citocromo c**, il quale verrà coinvolto in altri processi che si concluderanno con la morte della cellula. Le ROS hanno però diversi effetti positivi sul metabolismo e, nell'ospite dei mammiferi, possono proteggerlo dall'invasione dei **microbi**, probabilmente danneggiando il loro DNA mitocondriale. Per la loro capacità di danneggiare proteine, lipidi, DNA e RNA, esse sono coinvolte anche nel processo dell'invecchiamento, nella formazione delle cellule cancerogene, ma anche nella loro **necrosi** (una forma di morte cellulare incontrollata) e nell'*autofagia*](7).

(7) Tratto e modificato da: https://en.wikipedia.org/wiki/Reactive_oxygen_species

La grande importanza delle ROS nel nostro studio sull'**insulina** è comprensibile, in quanto la **produzione di insulina** e la **sua segnalazione** sono **processi redox sensibili** e la **compromissione della segnalazione fisiologica** da parte delle **ROS/RNS (Reactive Nitrogen Species)** è implicata nell'eziopatologia del *diabete* [Rochette L., Zeller M., Cottin Y., Vergely C. *Diabetes, oxidative stress and therapeutic strategies*. Biochim. Biophys. Acta. 2014; 1840:2709–2729. PMID: 24905298. DOI: 10.1016/j.bbagen.2014.05.017. <https://www.ncbi.nlm.nih.gov/pubmed/24905298>].

Per inciso, per una trattazione approfondita sulle ROS si consiglia lo studio del seguente articolo:

Egea J, Fabregat I, Frapart YM, et al. European contribution to the study of ROS: A summary of the findings and prospects for the future from the COST action BM1203 (EU-ROS). *Redox Biol.* 2017 Oct; 13: 94–162. PMID: 29054580. doi: 10.1016/j.redox.2017.05.007
<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC5458069>

Dagli articoli che invece riguardano la **ricerca mirata** sugli accoppiamenti dei termini “*Pseudomonas e Cystathionine Gamma-Synthase*” emergono principalmente **prove di forti legami** tra *Pseudomonas, Cystathionine Gamma-Synthase* e:

- 1) *Solfato di Idrogeno (H₂S)*;
- 2) *Metabolismo della Metionina*.

In conclusione, esistono **collegamenti indiretti** tra “*Insulin*” e “*Pseudomonas*” riconducibili a uno o più **percorsi metabolici** in cui è implicato **l'enzima CYSTATHIONINE GAMMA-SYNTHASE** (in particolare il metabolismo della *metionina* e quello dello *zolfo*).

1.15 CYTOCHROME C

Il risultato della ricerca BLAST identifica allineamenti significativi di basi con il CYTOCHROME C CLASS 1 dei batteri **Pseudomonas**. Questo risultato ha stimolato una **ricerca bibliografica specifica** sulle relazioni tra **Insulina**, **Pseudomonas** e **Cytochrome c**.

[Il **Cytochrome c** è una *emoproteina* (contenente *ferro*) capace di diffondere tra la membrana interna ed esterna dei *mitocondri*. Il gruppo *eme* è legato alla catena proteica tramite due legami covalenti con la *cisteina*. Il *Cytochrome c* è un componente essenziale della catena di trasporto degli elettroni dalla *Q-cytochrome c ossidoreduttasi* alla *cytochrome C ossidasi*. Il *Cytochrome c* ha anche un importante ruolo nel processo di *apoptosi* una forma di morte cellulare controllata finalizzata al ricambio cellulare o attivata in presenza di infezioni o danni del DNA. Esso è rilasciato dai mitocondri in risposta agli stimoli *pro-apoptotici*.](8). [Ad esempio, *Bax*, *BAD*, *Bak*, *Bok*, *etc.* (membri *pro-apoptotici* della famiglia genica *Bcl-2*), controllando la permeabilità della membrana mitocondriale, possono indurre il rilascio di **Cytochrome c** nel *citosol* (fluido intracellulare). Al contrario, *Bcl-2*, *Bcl-xL*, *Bcl-w*, *etc.* (membri *anti-apoptotici* della famiglia genica *Bcl-2*) possono invece inibirlo. La proteina *anti-apoptotica Bcl-2* impedisce il rilascio di **Cytochrome c** dai mitocondri e porta all'inibizione dell'apoptosi (probabilmente inibendo la proteina *Bax*). Al contrario, la proteina *Bax* consentendo un influsso di ioni attraverso la membrana mitocondriale esterna, favorisce il rilascio del **Cytochrome c** e incrementa l'apoptosi](9).

[Il **Cytochrome c1** è una proteina codificata dal **gene *CYCI*** e appartiene alla famiglia delle proteine del **Cytochrome c**. Il **Cytochrome c1** svolge un ruolo nel trasferimento di elettroni durante la fosforilazione ossidativa (il **Cytochrome c1**, insieme al *Cytocrome b*, trasferisce gli elettroni da CoQH_2 al *Cytochrome c*)](10).

[La **fosforilazione ossidativa** è un processo biochimico cellulare per la *produzione di ATP nei mitocondri*, fondamentale e ubiquitario. Si tratta della **fase finale della respirazione cellulare**. Essa è composta di due parti: la *catena di trasporto degli elettroni* e la *sintesi di ATP*](11).

(8) Tratto e modificato da: https://it.wikipedia.org/wiki/Citocromo_c

(9) Tratto e modificato da: <https://it.wikipedia.org/wiki/Bcl-2>

(10) Tratto e modificato da: https://en.wikipedia.org/wiki/Cytochrome_C1

(11) Tratto e modificato da: https://it.wikipedia.org/wiki/Fosforilazione_ossidativa

La ricerca bibliografica su tutti gli **articoli scientifici** pubblicati su **PubMed** in cui figurano insieme i termini “*Insulin*”, “*Pseudomonas*” e “*Cytochrome c*” **non ottiene risultati**

(<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+PSEUDOMONAS+CYTOCHROME+C>).

Questa constatazione fa ipotizzare possibili **relazioni indirette** tra *Insulin*, *Pseudomonas* e *Cytochrome C*.

Per indagare su queste **ipotetiche relazioni** si è proseguito facendo delle **ricerche mirate** sugli accoppiamenti dei termini “*Insulin e Cytochrome c (c1 e sue definizioni alternative)*” e “*Pseudomonas e Cytochrome c (c1 e sue definizioni alternative)*”.

Dagli articoli che riguardano la **ricerca mirata** sugli accoppiamenti dei termini “*Insulin e Cytochrome c (c1 e sue definizioni alternative)*” emergono principalmente **prove di forti legami** tra *Insulin*, *Cytochrome c* e:

- 1) *Apoptosi*;
- 2) *Stress ossidativo e le ROS (Reactive Oxygen Species)*;
- 3) *Diabete*;
- 4) *Obesità*.

Dagli articoli che riguardano la **ricerca mirata** sugli accoppiamenti dei termini “*Pseudomonas e Cytochrome c (c1 e sue definizioni alternative)*” emergono principalmente **prove di forti legami** tra *Pseudomonas*, *Cytochrome c* e:

- 1) *Apoptosi*;
- 2) *Stress ossidativo e le ROS (Reactive Oxygen Species)*;
- 3) *Fibrosi Cistica e Infezioni polmonari*.

Complessivamente, dai risultati della ricerca bibliografica presentata nelle pagine seguenti, si evidenzia, come fondamentale fattore comune tra **Insulina**, **Pseudomonas** e **Cytochrome c (c1 e su definizioni alternative)**, il ruolo determinante svolto dall'**attività mitocondriale**, soprattutto (ma non soltanto) in relazione allo **Stress Ossidativo**, alle **ROS** e all'**Apoptosi**.

1.16 HIGH-AFFINITY IRON TRANSPORTER

Il risultato della ricerca BLAST identifica allineamenti significativi di basi con un HIGH-AFFINITY IRON TRANSPORTER dei batteri **Pseudomonas**. Questo risultato ha stimolato una **ricerca bibliografica specifica** sulle relazioni tra **Insulina**, **Pseudomonas** e **Ferro**.

[Il **ferro** è presente abbondantemente nella Terra e rappresenta un micronutriente fondamentale per i vegetali e gli animali. Nel mondo animale, il ferro è inglobato nel *complesso eme*, un componente indispensabile delle proteine implicate nell'ossidazione (**reazioni redox**). Una tra le *reazioni redox* più importanti è la **respirazione cellulare**. Nel mondo inorganico il ferro si trova anche nelle azotasi e nelle idrogenasi (aggregati ferro-zolfo di vari enzimi). Altri *enzimi*, in cui il ferro è fondamentale, sono implicati in diverse funzioni, tra cui la *conversione del metano in metanolo* (*metano-monoossigenasi*) e la conversione del *ribosio in desossiribosio* (*ribonucleotide riduttasi*). Nel nostro organismo, per evitare un eccesso tossico, il ferro è *legato a proteine* che regolano il suo stato di ossidazione. La distribuzione degli *ioni ferro* nei mammiferi è quindi regolata in maniera molto precisa. Se, ad esempio, il corpo è soggetto a un'infezione, l'organismo **sottrae il ferro** rendendolo meno disponibile anche ai *batteri*. Alcune *proteine* che contengono ferro svolgono le funzioni principali di trasportare e conservare ossigeno e di trasferire elettroni. Il *ferro*, legato a **specifiche proteine di trasporto**, come l'*apoferritina* (che si trasforma in *ferritina*) e la *transferrina* (la principale *proteina* che lega il *ferro extracellulare* e lo trasporta nel sangue), è distribuito ai vari organi. Un tipo di ferro è idrosolubile (*ferro ferroso*) e più facilmente assorbibile dalle piante, mentre l'altro tipo è difficilmente solubile e assorbibile (*ferro ferrico*). Alla presenza di una concentrazione di ferro adeguata, i **trasportatori di ferro a bassa affinità** trasportano il *ferro ferroso* all'interno della cellula. Quando, invece, la concentrazione di ferro è scarsa, intervengono i **trasportatori di ferro ad alta affinità**. Esistono dei *recettori* (Recettori TfR1) per la *transferrina* ad **alta affinità** e dei *recettori* (Recettori TfR2) per la *transferrina* a **bassa affinità**. I Recettori TfR1 sono presenti negli *epatociti*, negli *enterociti* e nelle *cellule eritroidi*, mentre i Recettori TfR2 sono invece *ubiquitari*](12).

Per quello che riguarda le argomentazioni che si stanno sostenendo in questo lavoro, è necessario evidenziare lo **stretto rapporto** tra il **ferro** e i **citocromi**. Come già detto nel paragrafo 1.15 (pagina 42), i *citocromi* sono *proteine* che contengono *ferro* e sono deputate al trasporto degli elettroni, in particolare nel processo di fosforilazione ossidativa che porta alla produzione di diverse molecole di ATP. Il *ferro* e i *citocromi* sono quindi entrambi implicati nella **respirazione cellulare**.

(12) Tratto e modificato da: <https://it.wikipedia.org/wiki/Ferro>
<https://en.wikipedia.org/wiki/Iron>

Nel paragrafo 1.10 (pagina 35), è stato già evidenziato che, recentemente, per contrastare la virulenza di **Pseudomonas**, i ricercatori hanno sfruttato la sua forte avidità per il *ferro*. I *batteri* (compreso **Pseudomonas**) hanno sviluppato i *siderofori*, agenti che sequestrano il ferro ad **altissima affinità**. Gli *ioni di gallio* interagiscono con i processi cellulari in modo simile al *ferro (III)*. Quando gli *ioni di gallio* vengono scambiati per errore al posto del *ferro (III)* da batteri come *Pseudomonas*, gli ioni, **interagendo con l'attività dei citocromi**, interferiscono con la **respirazione cellulare** e causano la morte dei batteri. Ciò accade perché il *ferro è attivo redox*, consentendo il trasferimento di elettroni durante la respirazione, mentre il *gallio è inattivo redox*](13).

Questo argomento è possibile approfondirlo adeguatamente consultando il recentissimo lavoro di **Hijazi S., Visca P. e Frangipani E.** [*Gallium-Protoporphyrin IX Inhibits Pseudomonas aeruginosa Growth by Targeting Cytochromes*. Front Cell Infect Microbiol. 2017 Jan 26;7:12. doi: 10.3389/fcimb.2017.00012. eCollection 2017. PubMed PMID: 28184354; PubMed Central PMCID: PMC5266731. <https://www.ncbi.nlm.nih.gov/pubmed/28184354>].

La ricerca bibliografica su tutti gli **articoli scientifici** pubblicati su **PubMed** in cui figurano insieme i termini “*Insulin*”, “*Pseudomonas*” e “*Iron*” **non ottiene risultati** (<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+PSEUDOMONAS+IRON>).

Anche la ricerca bibliografica su tutti gli **articoli scientifici** pubblicati su **PubMed** in cui figurano insieme i termini “*Insulin*”, “*Burkholderia*” e “*Iron*” **non ottiene risultati** (<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+PSEUDOMONAS+IRON>).

Queste constatazioni fanno ipotizzare possibili **relazioni indirette** tra *Insulin*, *Pseudomonas* (o *Burkholderia*) e *Iron*.

Per indagare su queste **ipotetiche relazioni** si è proseguito facendo delle **ricerche mirate** sugli accoppiamenti dei termini “*Insulin e Iron*”, “*Pseudomonas e Iron*” e “*Burkholderia e Iron*”.

La quantità di articoli individuati dalle ricerche su questi accoppiamenti di termini è **enorme**. Per ridurre il campo d'indagine e rendere più agevole l'identificazione di relazioni significative (in riferimento agli scopi prefissi dal nostro lavoro) tra *Insulin* e *Iron*, tra *Pseudomonas* e *Iron*, e tra *Burkholderia* e *Iron*, sono stati inseriti altri **termini chiave** considerati importanti per delle **ricerche mirate**.

(13) Tratto e modificato da: <https://en.wikipedia.org/wiki/Pseudomonas>

Dagli articoli che riguardano i termini chiave, relativi agli accoppiamenti *Insulin/Iron*, *Pseudomonas/Iron* e *Burkholderia/Iron*, emergono principalmente **prove di forti legami** tra ognuno di questi accoppiamenti e i seguenti termini:

- 1) *Transporter*;
- 2) *Melioidosi*;
- 3) *Diabete*;
- 4) *Fibrosi Cistica*;
- 5) *Sistema Immunitario*;
- 6) *Apoptosi*;
- 7) *Cytochrome c*;
- 8) *Reactive Oxygen Species*;

Solo l'accoppiamento dei termini *Insulin* e *Iron*, **oltre ai risultati sopra esposti**, ha mostrato **prove di forti legami** con i seguenti termini:

- 1) *Obesità*;
- 2) *Insulite*.

In conclusione, per quel che riguarda i possibili legami tra *Insulina* e *Pseudomonas*, sembra essere **fondamentale anche il ruolo del ferro, dei suoi valori e del suo trasporto**.

1. 17 LA RICERCA RIGUARDANTE HELIGMOSOMOIDES POLYGYRUS

I risultati della ricerca BLAST (presentati a pagina 34) hanno stimolato un'opportuna ricerca bibliografica per studiare le **possibili relazioni** tra le **caratteristiche** della **sequenza originaria di DNA** (**63** basi della **Catena A dell'Insulina**) e le **caratteristiche** del nematoda **Heligmosomoides polygyrus** (o *Nematospiroides dubius*).

Dalla lettura degli articoli selezionati (presentati dalla pagina 157 alla 168) sono state riscontrate importanti relazioni, *dirette* o *indirette*, tra l'**insulina** e il nematoda **Heligmosomoides polygyrus**.

Queste importanti relazioni tra l'**insulina** e **Heligmosomoides polygyrus** avvalorano l'ipotesi che la **Sequenza n°1/1**, cioè la **nuova sequenza generata** rispettando fedelmente il **Trend n°1** (che è uno tra i possibili specifici "trend non manifesti" della *sequenza originaria*), abbia **forti relazioni** con le **caratteristiche** della **sequenza originaria** (nello specifico con la **Catena A dell'Insulina**).

In generale e molto sommariamente, tra le altre prove di minore importanza non citate, **esistono prove di forti legami** (anche se il numero di articoli, in alcuni casi, è scarso) tra *Insulina*, *Heligmosomoides polygyrus* e:

- 1) *Sistema Immunitario*;
- 2) *Apoptosi*;
- 3) *Infezioni Polmonari*;
- 4) *Diabete*;
- 5) *Obesità*;
- 6) *Insulite*.

Sono state fatte anche delle **ricerche mirate** sulle relazioni esistenti tra *Heligmosomoides polygyrus* e i singoli **termini rilevanti** evidenziati ai punti 1, 2, 3, 4, 5 e 6.

In un articolo (**non consultabile online**) è stata trovata una corrispondenza tra i termini *Heligmosomoides polygyrus* e *Pseudomonas*.

In linea generale, i **risultati conclusivi** di queste ricerche, provano che l'infezione da *Heligmosomoides polygyrus* può ridurre, attraverso una forte risposta immunitaria dell'ospite, il *diabete di tipo 1*, la tendenza a prendere eccessivo peso (*obesità*), molte *infezioni polmonari* (ad eccezione della polmonite da pneumococco causata da *Streptococcus pneumoniae*) e l'*apoptosi*.

RISULTATI DELLA RICERCA BIBLIOGRAFICA SU PSEUDOMONAS E SU HELIGMOSOMOIDES POLYGYRUS

1.18 CONCLUSIONI CAPITOLO I° (PARTE PRIMA)

Nell'insieme, rispetto ai batteri **Pseudomonas**, dalla ricerca bibliografica emergono numerose e importanti relazioni tra diverse specie di batteri **Pseudomonas** (identificati dalla ricerca BLAST per gli allineamenti significativi tra la **Sequenza n°1/1** e alcune loro *basi di DNA* inerenti la *Cystathionine Gamma-Synthase*, i *Citocromi C classe 1* e un *High-Affinity Iron Transporter*) e l'**insulina**, il **diabete mellito**, la **melioidosi**, l'**obesità**, la **fibrosi cistica**, diversi tipi d'**infezioni** (soprattutto **polmonari**), l'**otite esterna maligna**, l'**endocardite**, il **sistema immunitario**, l'**apoptosi** e i **valori del ferro** e il suo **trasporto**.

Anche riguardo, il nemoda **Heligmosomoides polygyrus**, la ricerca bibliografica evidenzia numerose e importanti relazioni tra questo *parassita intestinale* (identificato dalla ricerca BLAST per gli allineamenti significativi tra la **Sequenza n°1/1** e alcune sue *basi di DNA*) e l'**insulina**, il **sistema immunitario**, l'**apoptosi**, il **diabete di tipo I**, l'**obesità** e l'**insulite**.

Non è stato invece possibile indagare sulle relazioni tra *Pseudomonas* e *Heligmosomoides polygyrus*, perché in rete è stato rintracciato un solo articolo, peraltro molto datato e non consultabile online.

Sarebbero augurabili futuri studi specifici, soprattutto perché alcune caratteristiche inerenti *Pseudomonas* appaiono **speculari** e **opposte** a quelle manifestate da *Heligmosomoides polygyrus*; infatti, correlata alla sua infezione, si riscontra nell'ospite una riduzione della gravità del **diabete di tipo 1**, dell'**obesità**, dell'**insulite** e dell'**apoptosi** (mentre, al contrario, l'infezione da *Pseudomonas* è correlata al loro incremento di gravità).

Prima di passare alla seconda parte di questo capitolo, si sottolinea l'importanza evidenziata dalla ricerca bibliografica riguardo alla **respirazione cellulare**, le **ROS** (*Reactive Oxygen Species*), i **valori del ferro** e il suo **trasporto**, i quali (insieme o separatamente) sembrano rappresentare *un punto di contatto importante* tra i diversi aspetti dei fenomeni e delle patologie considerate in questa prima parte del capitolo.

L'**ipotesi** che s'intendeva avvalorare, cioè che la **nuova sequenza** (generata rispettando fedelmente uno tra i possibili specifici "trend non manifesti" della **sequenza originaria**) avesse forti relazioni con le *caratteristiche* della **sequenza originaria** (nello specifico con la **Catena A dell'insulina**) sembra essere sostenibile e verosimile.

La ricerca bibliografica sembra confermare che l'**insulina** (e, quindi, anche la **Catena A dell'insulina**) è in vari modi *molto implicata con alcune caratteristiche* sia dei batteri **Pseudomonas**, sia del nemoda **Heligmosomoides polygyrus** (*organismi entrambi costituiti da basi di DNA della nuova sequenza generata*).

In conclusione, l'analisi (attraverso la **T.T.E.S.**) della *sequenza originaria* basata su uno dei suoi "trend non manifesti" (*il Trend n°1*), la creazione di una *nuova sequenza* di DNA (la **Sequenza n°1/1**) a partire dal **Trend n°1** della *sequenza originaria* e la *congruenza* con i dati ottenuti dall'*approfondimento bibliografico* aprono prospettive totalmente inesplorate riguardo all'ambito della ricerca genetica e delle sue innumerevoli applicazioni.

1.19 LE PROSPETTIVE DI RICERCA DEI PROSSIMI CAPITOLI

La ricerca BLAST della *nuova sequenza di DNA* analizzata in questo capitolo, generata rispettando fedelmente uno tra i possibili "trend non manifesti" (cioè *il Trend n°1*), ha dato anche altri risultati di *allineamenti significativi di basi* oltre i batteri **Pseudomonas** e il nemoda **Heligmosomoides polygyrus**.

Nelle altre parti di questo capitolo, lo stesso tipo di ricerca bibliografica sarà quindi eseguita su tutti gli altri *organismi* per i quali si sono evidenziati, dalla ricerca BLAST, allineamenti significati alla **Sequenza n°1/1**.

Dal nostro punto di vista, ogni *organismo* per il quale è stato riscontrato un allineamento significativo con la **nuova sequenza generata (Sequenza n°1/1)** dovrebbe, a ragion veduta, essere in qualche modo legato (direttamente o indirettamente) alla **sequenza originaria** (nello specifico alla **Catena A dell'Insulina**), agli *organismi* identificati con la ricerca Blast eseguita sulla **Catena A dell'Insulina** e, in parte, anche agli altri *organismi* per i quali saranno riscontrati allineamenti significativi con **altre sequenze generate**.

Indagare anche su questi legami sarà un *obiettivo* dei prossimi capitoli di questa *nuova e affascinante prospettiva d'indagine genetica*.

Un *altro obiettivo*, molto più arduo, lungo e impegnativo, sarà quello di analizzare i risultati di *tutti gli allineamenti significativi* ottenuti da *tutte le possibili sequenze generate* rispettando fedelmente *tutti i possibili specifici* "trend non manifesti" della *sequenza originaria* (nello specifico la **Catena A dell'Insulina**).

Bibliografia scientifica selezionata

BIBLIOGRAFIA RIGUARDANTE PSEUDOMONAS

1.20 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA E PSEUDOMONAS (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “*Insulina*” e “*Pseudomonas*”.

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+PSEUDOMONAS>

Sort by: Most Recent - Search results = Items: 147

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi**:

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1.21 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA E PSEUDOMONAS (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle migliori corrispondenze tra i termini “*Insulina*” e “*Pseudomonas*” (**Attenzione:** gli articoli già citati nella ricerca precedente non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+PSEUDOMONAS>

Sort by: Best Match - Search results = Item: 134

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

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1. 22 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, PSEUDOMONAS E DIABETE (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “*Insulina*”, “*Pseudomonas*” e “*Diabete*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+PSEUDOMONAS+DIABETES>

Sort by: Most Recent - Search results = Items: 49

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Snarski E, Milczarczyk A, Halaburda K, Torosian T, Paluszewska M, Urbanowska E, Król M, Boguradzki P, Jedynasty K, Franek E, Wiktor-Jedrzejczak W. *Immunoablation and autologous hematopoietic stem cell transplantation in the treatment of new-onset type 1 diabetes mellitus: long-term observations.* Bone Marrow Transplant. 2016 Mar;51(3):398-402. doi: 10.1038/bmt.2015.294. Epub 2015 Dec 7. PubMed PMID: 26642342.
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1.23 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA E BURKHOLDERIA (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “*Insulina*” e “*Burkholderia*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+BURKHOLDERIA>

Sort by: **Most Recent** - *Search results = Items: 23*

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1.24 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA E CYSTATHIONINE GAMMA-SYNTASE

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) dei termini “*Insulina*” e “*Cystathionine Gamma-Synthase*”.

<https://www.ncbi.nlm.nih.gov/pmc/?term=INSULIN+CYSTATHIONINE+GAMMA-SYNTASE>

Search results = Items: 16

Dai risultati della ricerca sono stati selezionati gli **articoli ritenuti più significativi**:

Hak Joo Lee, Meenalakshmi M. Mariappan, Denis Feliars, Rita C. Cavaglieri, Kavithalakshmi Sataranatarajan, Hanna E. Abboud, Goutam Ghosh Choudhury, Balakuntalam S. Kasinath. *Hydrogen Sulfide Inhibits High Glucose-induced Matrix Protein Synthesis by Activating AMP-activated Protein Kinase in Renal Epithelial Cells.* J Biol Chem. 2012 Feb 10; 287(7): 4451–4461. Published online 2011 Dec 9. doi: 10.1074/jbc.M111.278325 PMID: PMC3281646. <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3281646/>

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Shaimaa S. El-Sayed, Mohamed NM. Zakaria, Rasha H. Abdel-Ghany, Abdel A. Abdel-Rahman. *Cystathionine- γ lyase-derived hydrogen sulfide mediates the cardiovascular protective effects of moxonidine in diabetic rats.* Eur J Pharmacol. Author manuscript; available in PMC 2017 Jul 15. Published in final edited form as: Eur J Pharmacol. 2016 Jul 15; 783: 73–84. Published online 2016 Apr 29. doi: 10.1016/j.ejphar.2016.04.054 PMID: PMC4893977. <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4893977>

Fuqiang Yin, Agnieszka Pajak, Ralph Chapman, Andrew Sharpe, Shangzhi Huang, Frédéric Marsolais. *Analysis of common bean expressed sequence tags identifies sulfur metabolic pathways active in seed and sulfur-rich proteins highly expressed in the absence of phaseolin and major lectins.* BMC Genomics. 2011; 12: 268. Published online 2011 May 26. doi: 10.1186/1471-2164-12-268 PMID: PMC3115882. <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3115882>

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Elosie Y Streeter, Emilio Badoer, Owen L Woodman, Joanne L Hart. *Effect of type 1 diabetes on the production and vasoactivity of hydrogen sulfide in rat middle cerebral arteries.* Physiol Rep. 2013 Oct; 1(5): e00111. Published online 2013 Oct 20. doi: 10.1002/phy2.111 PMID: PMC3841046. <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3841046>

Heidi M. Blank, Shefali Gajjar, Andrey Belyanin, Michael Polymenis. *Sulfur Metabolism Actively Promotes Initiation of Cell Division in Yeast.* PLoS One. 2009; 4(11): e8018. Published online 2009 Nov 24. doi: 10.1371/journal.pone.0008018 PMID: PMC2776973.
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Matthew Whiteman, Philip K Moore. *Hydrogen sulfide and the vasculature: a novel vasculoprotective entity and regulator of nitric oxide bioavailability?* J Cell Mol Med. 2009 Mar; 13(3): 488–507. Published online 2009 Mar 24. doi: 10.1111/j.1582-4934.2009.00645.x PMID: PMC3822510. <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3822510>

1.25 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, HOMOCYSTEINE E HYDROGEN SULFIDE SYSTEM

Per lo studio di questi argomenti si consultino i risultati della ricerca ottenuta usando il **Link** mostrato qui di seguito:

https://www.ncbi.nlm.nih.gov/pubmed?db=pubmed&cmd=link&linkname=pubmed_pubmed&uid=10470367

Search results = Items: 91

Concludo con **due articoli** e con la tesi di dottorato di **Ling Zhang** (2012), entrambi interessati al rapporto tra *Insulina*, metabolismo della *Cisteina* e ruolo del *Solfato di Idrogeno*:

Kimura H. *Hydrogen sulfide: its production, release and functions.* Amino Acids. 2011 Jun;41(1):113-21. PMID: 20191298. DOI: 10.1007/s00726-010-0510-x.
<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+HOMOCYSTEINE+HYDROGEN+SULFIDE+SYSTEM>

Yukiko Kaneko, Yuka Kimura, Hideo Kimura, Ichiro Niki. *l-Cysteine Inhibits Insulin Release From the Pancreatic β -Cell. Possible Involvement of Metabolic Production of Hydrogen Sulfide, a Novel Gasotransmitter.* Diabetes May 2006, 55 (5) 1391-1397. PMID: 16644696. DOI: 10.2337/db05-1082. <https://www.ncbi.nlm.nih.gov/pubmed/16644696>

L. Zhang. *Cystathionine gamma-lyase/hydrogen sulfide system and glucose homeostasis.* Lakehead University. 12-Feb-2013. <http://knowledgecommons.lakeheadu.ca/handle/2453/443>

1.26 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS E CYSTATHIONINE GAMMA-SYNTHASE

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica su tutti gli **articoli scientifici** pubblicati su **PubMed** in cui figurano insieme i termini “*Pseudomonas*” e “*Cystathionine Gamma-Synthase*”.

<https://www.ncbi.nlm.nih.gov/pubmed/?term=Pseudomonas+CYSTATHIONINE+GAMMA-SYNTHASE>

Search results = Items: 6

Motoshima H, Inagaki K, Kumasaka T, Furuichi M, Inoue H, Tamura T, Esaki N, Soda K, Tanaka N, Yamamoto M, Tanaka H. *Crystal structure of the pyridoxal 5'-phosphate dependent L-methionine gamma-lyase from Pseudomonas putida.* J Biochem. 2000 Sep;128(3):349-54. PubMed PMID: 10965031. <https://www.ncbi.nlm.nih.gov/pubmed/10965031>

Vermeij P, Kertesz MA. *Pathways of assimilative sulfur metabolism in Pseudomonas putida.* J Bacteriol. 1999 Sep;181(18):5833-7. PubMed PMID: 10482527; PubMed Central PMCID: PMC94106. <https://www.ncbi.nlm.nih.gov/pubmed/10482527>

Taté R, Riccio A, Caputo E, Iaccarino M, Patriarca EJ. *The Rhizobium etli metZ gene is essential for methionine biosynthesis and nodulation of Phaseolus vulgaris.* Mol Plant Microbe Interact. 1999 Jan;12(1):24-34. PubMed PMID: 9885190. <https://www.ncbi.nlm.nih.gov/pubmed/9885190>

Inoue H, Inagaki K, Sugimoto M, Esaki N, Soda K, Tanaka H. *Structural analysis of the L-methionine gamma-lyase gene from Pseudomonas putida.* J Biochem. 1995 May;117(5):1120-5. PubMed PMID: 8586629. <https://www.ncbi.nlm.nih.gov/pubmed/8586629>

Fogolino M, Borne F, Bally M, Ball G, Patte JC. *A direct sulfhydrylation pathway is used for methionine biosynthesis in Pseudomonas aeruginosa.* Microbiology. 1995 Feb;141 (Pt 2):431-9. PubMed PMID: 7704274. <https://www.ncbi.nlm.nih.gov/pubmed/7704274>

Nagasawa T, Kanzaki H, Yamada H. *Cystathionine gamma-lyase of Streptomyces phaeochromogenes. The occurrence of cystathionine gamma-lyase in filamentous bacteria and its purification and characterization.* J Biol Chem. 1984 Aug 25;259(16):10393-403. PubMed PMID: 6432781. <https://www.ncbi.nlm.nih.gov/pubmed/6432781>

Prima di concludere, presento altri **due interessanti articoli** sull'argomento:

G L Andersen, G A Beattie and S E Lindow. *Molecular Characterization and Sequence of a Methionine Biosynthetic Locus from Pseudomonas syringae.* Journal of Bacteriology, 1998. Vol. 180 Iss. 17 (1998) p. 4497 - 4507. PMCID: PMC107460. <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC107460/>

Hacham Y., Gophna U. e Amir R. *In vivo analysis of various substrates utilized by cystathionine gamma-synthase and O-acetylhomoserine sulfhydrylase in methionine biosynthesis.* Molecular Biology and Evolution, Volume 20, Issue 9, 1 September 2003, Pages 1513–1520. PMID: 12832650. <https://doi.org/10.1093/molbev/msg169>

1.27 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA E CYTOCHROME C (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Insulina*” e “*Cytochrome c*”.

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+CYTOCHROME+C>

Sort by: Most Recent - Search results = Items: 745

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi**:

Beyfuss K, Hood DA. *A systematic review of p53 regulation of oxidative stress in skeletal muscle.* Redox Rep. 2018 Jan 3:1-18. doi: 10.1080/13510002.2017.1416773. [Epub ahead of print] PubMed PMID: 29298131. <https://www.ncbi.nlm.nih.gov/pubmed/29298131>

Yuan F, Woollard JR, Jordan KL, Lerman A, Lerman LO, Eirin A. *Mitochondrial Targeted Peptides Preserve Mitochondrial Organization and Decrease Reversible Myocardial Changes in Early Swine Metabolic Syndrome.* Cardiovasc Res. 2017 Dec 18. doi: 10.1093/cvr/cvx245. [Epub ahead of print] PubMed PMID: 29267873. <https://www.ncbi.nlm.nih.gov/pubmed/29267873>

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Park HS, Cho HS, Kim TW. *Physical exercise promotes memory capability by enhancing hippocampal mitochondrial functions and inhibiting apoptosis in obesity-induced insulin resistance by high fat diet.* Metab Brain Dis. 2017 Nov 29. doi: 10.1007/s11011-017-0160-8. [Epub ahead of print] PubMed PMID: 29185193. <https://www.ncbi.nlm.nih.gov/pubmed/29185193>

Feng CC, Pandey S, Lin CY, Shen CY, Chang RL, Chang TT, Chen RJ, Viswanadha VP, Lin YM, Huang CY. *Cardiac apoptosis induced under high glucose condition involves activation of IGF2R signaling in H9c2 cardiomyoblasts and streptozotocin-induced diabetic rat hearts.* Biomed Pharmacother. 2017 Nov 6;97:880-885. doi: 10.1016/j.biopha.2017.11.020. [Epub ahead of print] PubMed PMID: 29136764. <https://www.ncbi.nlm.nih.gov/pubmed/29136764>

Zhang Y, Wang M, Dong H, Yu X, Zhang J. *Anti-hypoglycemic and hepatocyte-protective effects of hyperoside from Zanthoxylum bungeanum leaves in mice with high-carbohydrate/high-fat diet and alloxan-induced diabetes.* Int J Mol Med. 2018 Jan;41(1):77-86. doi: 10.3892/ijmm.2017.3211. Epub 2017 Oct 25. PubMed PMID: 29115390. <https://www.ncbi.nlm.nih.gov/pubmed/29115390>

Liu CM, Ma JQ, Sun JM, Feng ZJ, Cheng C, Yang W, Jiang H. *Association of changes in ER stress-mediated signaling pathway with lead-induced insulin resistance and apoptosis in rats and their prevention by A-type dimeric epigallocatechin-3-gallate.* Food Chem Toxicol. 2017 Dec;110:325-332. doi: 10.1016/j.fct.2017.10.040. Epub 2017 Oct 27. PubMed PMID: 29107025. <https://www.ncbi.nlm.nih.gov/pubmed/29107025>

Xie X, Sinha S, Yi Z, Langlais PR, Madan M, Bowen BP, Willis W, Meyer C. *Role of adipocyte mitochondria in inflammation, lipemia and insulin sensitivity in humans: effects of pioglitazone treatment.* Int J Obes (Lond). 2017 Aug 14. doi: 10.1038/ijo.2017.192. [Epub ahead of print] PubMed PMID: 29087390. <https://www.ncbi.nlm.nih.gov/pubmed/29087390>

Onyango AN. *The Contribution of Singlet Oxygen to Insulin Resistance.* Oxid Med Cell Longev. 2017;2017:8765972. doi: 10.1155/2017/8765972. Epub 2017 Sep 7. Review. PubMed PMID: 29081894; PubMed Central PMCID: PMC5610878. <https://www.ncbi.nlm.nih.gov/pubmed/29081894>

Jiménez-Maldonado A, Ying Z, Byun HR, Gomez-Pinilla F. *Short-term fructose ingestion affects the brain independently from establishment of metabolic syndrome.* Biochim Biophys Acta. 2018 Jan;1864(1):24-33. doi: 10.1016/j.bbadis.2017.10.012. Epub 2017 Oct 7. PubMed PMID: 29017895; PubMed Central PMCID: PMC5705281. <https://www.ncbi.nlm.nih.gov/pubmed/29017895>

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Aghanoori MR, Smith DR, Roy Chowdhury S, Sabbir MG, Calcutt NA, Fernyhough P. *Insulin prevents aberrant mitochondrial phenotype in sensory neurons of type 1 diabetic rats.* Exp Neurol. 2017 Nov;297:148-157. doi: 10.1016/j.expneurol.2017.08.005. Epub 2017 Aug 10. PubMed PMID: 28803751; PubMed Central PMCID: PMC5612919. <https://www.ncbi.nlm.nih.gov/pubmed/28803751>

Othman AI, El-Sawi MR, El-Missiry MA, Abukhalil MH. *Epigallocatechin-3-gallate protects against diabetic cardiomyopathy through modulating the cardiometabolic risk factors, oxidative stress, inflammation, cell death and fibrosis in streptozotocin-nicotinamide-induced diabetic rats.* Biomed Pharmacother. 2017 Oct;94:362-373. doi: 10.1016/j.biopha.2017.07.129. Epub 2017 Aug 1. PubMed PMID: 28772214. <https://www.ncbi.nlm.nih.gov/pubmed/28772214>

Silvander JSG, Kvarnström SM, Kumari-Ilieva A, Shrestha A, Alam CM, Toivola DM. *Keratins regulate β -cell mitochondrial morphology, motility, and homeostasis.* FASEB J. 2017 Oct;31(10):4578-4587. doi: 10.1096/fj.201700095R. Epub 2017 Jun 30. PubMed PMID: 28666985. <https://www.ncbi.nlm.nih.gov/pubmed/28666985>

Wilson DF, Cember ATJ, Matschinsky FM. *The thermodynamic basis of glucose-stimulated insulin release: a model of the core mechanism.* Physiol Rep. 2017 Jun;5(12). pii: e13327. doi: 10.14814/phy2.13327. PubMed PMID: 28655753; PubMed Central PMCID: PMC5492210. <https://www.ncbi.nlm.nih.gov/pubmed/28655753>

Wei XB, Guo L, Liu Y, Zhou SR, Liu Y, Dou X, Du SY, Ding M, Peng WQ, Qian SW, Huang HY, Tang QQ. *Synthesis of cytochrome c oxidase I (SCO1) inhibits insulin sensitivity by decreasing copper levels in adipocytes.* Biochem Biophys Res Commun. 2017 Sep 23;491(3):814-820. doi: 10.1016/j.bbrc.2017.06.124. Epub 2017 Jun 21. PubMed PMID: 28647369.
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Martín-Montañez E, Millon C, Boraldi F, Garcia-Guirado F, Pedraza C, Lara E, Santin LJ, Pavia J, Garcia-Fernandez M. *IGF-II promotes neuroprotection and neuroplasticity recovery in a long-lasting model of oxidative damage induced by glucocorticoids.* Redox Biol. 2017 Oct;13:69-81. doi: 10.1016/j.redox.2017.05.012. Epub 2017 May 26. PubMed PMID: 28575743; PubMed Central PMCID: PMC5454142. <https://www.ncbi.nlm.nih.gov/pubmed/28575743>

Candeias E, Sebastião I, Cardoso S, Carvalho C, Santos MS, Oliveira CR, Moreira PI, Duarte AI. *Brain GLP-1/IGF-1 Signaling and Autophagy Mediate Exendin-4 Protection Against Apoptosis in Type 2 Diabetic Rats.* Mol Neurobiol. 2017 Jun 2. doi: 10.1007/s12035-017-0622-3. [Epub ahead of print] PubMed PMID: 28573460. <https://www.ncbi.nlm.nih.gov/pubmed/28573460>

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Zhao B, Zheng Z. *Insulin Growth Factor 1 Protects Neural Stem Cells Against Apoptosis Induced by Hypoxia Through Akt/Mitogen-Activated Protein Kinase/Extracellular Signal-Regulated Kinase (Akt/MAPK/ERK) Pathway in Hypoxia-Ishchemic Encephalopathy.* Med Sci Monit. 2017 Apr 19;23:1872-1879. PubMed PMID: 28420864; PubMed Central PMCID: PMC5405785.
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<https://www.ncbi.nlm.nih.gov/pubmed/28122051>

1.28 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA E CYTOCHROME C (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle migliori corrispondenze tra i termini “*Insulina*” e “*Cytochrome c*” (**Attenzione:** gli articoli già citati nella ricerca precedente non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+CYTOCHROME+C>

Sort by: **Best Match** - Search results = Item: 626

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi**:

Rountree AM, Neal AS, Lisowski M, Rizzo N, Radtke J, White S, Luciani DS, Kim F, Hampe CS, Sweet IR. *Control of insulin secretion by cytochrome C and calcium signaling in islets with impaired metabolism.* J Biol Chem. 2014 Jul 4;289(27):19110-9. doi: 10.1074/jbc.M114.556050. Epub 2014 May 19. PubMed PMID: 24841202; PubMed Central PMCID: PMC4081948.
<https://www.ncbi.nlm.nih.gov/pubmed/24841202>

Sanderson TH, Mahapatra G, Pecina P, Ji Q, Yu K, Sinkler C, Varughese A, Kumar R, Bukowski MJ, Tousignant RN, Salomon AR, Lee I, Hüttemann M. *Cytochrome C is tyrosine 97 phosphorylated by neuroprotective insulin treatment.* PLoS One. 2013 Nov 5;8(11):e78627. doi: 10.1371/journal.pone.0078627. eCollection 2013. PubMed PMID: 24223835; PubMed Central PMCID: PMC3818486. <https://www.ncbi.nlm.nih.gov/pubmed/24223835>

Jung SR, Kuok IT, Couron D, Rizzo N, Margineantu DH, Hockenbery DM, Kim F, Sweet IR. *Reduced cytochrome C is an essential regulator of sustained insulin secretion by pancreatic islets.* J Biol Chem. 2011 May 20;286(20):17422-34. doi: 10.1074/jbc.M110.202820. Epub 2011 Mar 10. PubMed PMID: 21393241; PubMed Central PMCID: PMC3093816.
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P. Gaignard, M. Menezes, M. Schiff, A. Bayot, M. Rak, Ogier de Baulny, C, Su, M. Gilleron, A. Lombes, H. Abida, A. Tzagoloff, L. Riley, S. T. Cooper, K. Mina, P. Sivadorai, M, R. Davis, R. J. N. Allcock, N. Kresoje, N. G. Laing, D. R. Thorburn, A. Slama, J. Christodoulou and P. Rustin. *Mutations in CYC1, Encoding Cytochrome c1 Subunit of Respiratory Chain Complex III, Cause Insulin-Responsive Hyperglycemia.* AJHG, 8 August 2013, Volume 93, Issue 2, p 384–389. doi: 10.1016/j.ajhg.2013.06.015. Epub 2013 Aug 1. PubMed PMID: 23910460; PubMed C. PMCID: PMC3738829. <https://www.ncbi.nlm.nih.gov/pubmed/23910460>

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<https://www.ncbi.nlm.nih.gov/pubmed/24841383>

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Sweet IR, Gilbert M, Jensen R, Sabek O, Fraga DW, Gaber AO, Reems J. *Glucose stimulation of cytochrome C reduction and oxygen consumption as assessment of human islet quality. Transplantation.* 2005 Oct 27;80(8):1003-11. PubMed PMID: 16278578.

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Wu H, Xu G, Liao Y, Ren H, Fan J, Sun Z, Zhang M. *Supplementation with antioxidants attenuates transient worsening of retinopathy in diabetes caused by acute intensive insulin therapy.* Graefes Arch Clin Exp Ophthalmol. 2012 Oct;250(10):1453-8. doi: 10.1007/s00417-012-2079-4. Epub 2012 Jun 14. PubMed PMID: 22695936. <https://www.ncbi.nlm.nih.gov/pubmed/22695936>

Li Y, Higashi Y, Itabe H, Song YH, Du J, Delafontaine P. *Insulin-like growth factor-1 receptor activation inhibits oxidized LDL-induced cytochrome C release and apoptosis via the phosphatidylinositol 3 kinase/Akt signaling pathway.* Arterioscler Thromb Vasc Biol. 2003 Dec;23(12):2178-84. Epub 2003 Oct 9. PubMed PMID: 14551153.

<https://www.ncbi.nlm.nih.gov/pubmed/14551153>

1.29 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA E CYTOCHROME C1 (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Insulina*” e “*Cytochrome c1*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+CYTOCHROME+C1>

Sort by: Most Recent - Search results = Item: 12

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Alecu I, Othman A, Penno A, Saied EM, Arenz C, von Eckardstein A, Hornemann T. *Cytotoxic 1-deoxysphingolipids are metabolized by a cytochrome P450-dependent pathway.* J Lipid Res. 2017 Jan;58(1):60-71. doi: 10.1194/jlr.M072421. Epub 2016 Nov 21. PubMed PMID: 27872144; PubMed Central PMCID: PMC5234722.

<https://www.ncbi.nlm.nih.gov/pubmed/27872144>

Sugizaki T, Watanabe M, Horai Y, Kaneko-Iwasaki N, Arita E, Miyazaki T, Morimoto K, Honda A, Irie J, Itoh H. *The Niemann-Pick C1 like 1 (NPC1L1) inhibitor ezetimibe improves metabolic disease via decreased liver X receptor (LXR) activity in liver of obese male mice.* Endocrinology. 2014 Aug;155(8):2810-9. doi: 10.1210/en.2013-2143. Epub 2014 Apr 28. PubMed PMID: 24773344. <https://www.ncbi.nlm.nih.gov/pubmed/24773344>

Shertzer HG, Krishan M, Genter MB. *Dietary whey protein stimulates mitochondrial activity and decreases oxidative stress in mouse female brain.* Neurosci Lett. 2013 Aug 26;548:159-64. doi: 10.1016/j.neulet.2013.05.061. Epub 2013 Jun 6. PubMed PMID: 23748211; PubMed Central PMCID: PMC3749878. <https://www.ncbi.nlm.nih.gov/pubmed/23748211>

Sharma TS, Jacobson DL, Anderson L, Gerschenson M, Van Dyke RB, McFarland EJ, Miller TL; Pediatric HIV/AIDS Cohort Study (PHACS). *Short communication: The relationship between mitochondrial dysfunction and insulin resistance in HIV-infected children receiving antiretroviral therapy.* AIDS Res Hum Retroviruses. 2013 Sep;29(9):1211-7. PubMed PMID: 23742635; PubMed Central PMCID: PMC3749716. <https://www.ncbi.nlm.nih.gov/pubmed/23742635>

M. H. Holmström, R. Z. Tom, M. Björnholm, P. M. Garcia-Roves and J. R. Zierath. *Effect of leptin treatment on mitochondrial function in obese leptin-deficient ob/ob mice.* Metabolism. 2013 Sep; 62(9): 1258-67. doi: 10.1016/j.metabol.2013.04.001. Epub 2013 May 8. PubMed PMID: 23664724. <https://www.ncbi.nlm.nih.gov/pubmed/23664724>

A. Gesing, M. M. Masternak, F. Wang, M. Karbownik-Lewinska and A. Bartke. *Deletion of growth hormone receptor gene but not visceral fat removal decreases expression of apoptosis-related genes in the kidney-potential mechanism of lifespan extension.* Age (Dordr). 2012 Apr; 34(2): 295-304. doi: 10.1007/s11357-011-9232-6. Epub 2011 Mar 23. PubMed PMID: 21431351; PubMed Central PMCID: PMC3312636. <https://www.ncbi.nlm.nih.gov/pubmed/21431351>

E. J. Wilson and W. C. McMurray. *Effects of hormones on the maintenance and mitochondrial functions of rat hepatocytes cultured in serum-free medium.* Can J Biochem Cell Biol., 1983 Jul; 61(7):636-43. PMID: 6354397. <https://www.ncbi.nlm.nih.gov/pubmed/6354397>

Y. Ogawa, K. Matsumoto and S. Ofuji. *Changes in adenine nucleotide and mitochondrial metabolism of the kidney of burned rats and their relation to insulin.* J Lab Clin Med. 1977 Sep; 90(3): 457-65. PMID: 894101. <https://www.ncbi.nlm.nih.gov/pubmed/894101/>

1.30 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA E LE DENOMINAZIONI ALTERNATIVE DEL CYTOCHROME C1 (UNA SELEZIONE DELLE PIU' RECENTI E MIGLIORI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* o *migliori* corrispondenze tra i termini “*Insulina*” e le denominazioni alternative del “**Cytochrome c1**”: “*Cytochrome c-1*”, “*Complex III Subunit IV*” e “*Complex III Subunit 4*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

1) Cytochrome c-1:

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+CYTOCHROME+C-1>

Sort by: Best Match - Search results = Item: 13

Anastasio N, Tarailo-Graovac M, Al-Khalifah R, Legault L, Drogemoller B, Ross CJ, Wasserman WW, van Karnebeek C, Buhas D. *Mitochondrial Complex III Deficiency with Ketoacidosis and Hyperglycemia Mimicking Neonatal Diabetes*. JIMD Rep. 2017;31:57-62. doi: 10.1007/8904_2016_557. Epub 2016 Apr 14. PubMed PMID: 27074787; PubMed Central PMCID: PMC5388639. <https://www.ncbi.nlm.nih.gov/pubmed/27074787>

2) Complex III Subunit IV:

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+COMPLEX+III+SUBUNIT+IV>

Sort by: Most Recent - Search results = Item: 18

Rowley TJ 4th, Bitner BF, Ray JD, Lathen DR, Smithson AT, Dallon BW, Plowman CJ, Bikman BT, Hansen JM, Dorenkott MR, Goodrich KM, Ye L, O'Keefe SF, Neilson AP, Tessem JS. *Monomeric cocoa catechins enhance β -cell function by increasing mitochondrial respiration*. J Nutr Biochem. 2017 Nov;49:30-41. doi: 10.1016/j.jnutbio.2017.07.015. Epub 2017 Jul 27. PubMed PMID: 28863367. <https://www.ncbi.nlm.nih.gov/pubmed/28863367>

Shiba S, Ikeda K, Horie-Inoue K, Nakayama A, Tanaka T, Inoue S. *Deficiency of COX7RP, a mitochondrial supercomplex assembly promoting factor, lowers blood glucose level in mice*. Sci Rep. 2017 Aug 8;7(1):7606. doi: 10.1038/s41598-017-08081-z. PubMed PMID: 28790391; PubMed Central PMCID: PMC5548899. <https://www.ncbi.nlm.nih.gov/pubmed/28790391>

Stefano GB, Challenger S, Kream RM. *Hyperglycemia-associated alterations in cellular signaling and dysregulated mitochondrial bioenergetics in human metabolic disorders*. Eur J Nutr. 2016 Dec;55(8):2339-2345. Epub 2016 Apr 15. Review. PubMed PMID: 27084094; PubMed Central PMCID: PMC5122622. <https://www.ncbi.nlm.nih.gov/pubmed/27084094>

Amengual-Cladera E, Capllonch-Amer G, Lladó I, Gianotti M, Proenza AM. *Proteomic study of periovarian adipose tissue in 17 β -estradiol-treated and untreated ovariectomized rats*. Biochem Cell Biol. 2016 Apr;94(2):167-75. doi: 10.1139/bcb-2015-0077. Epub 2016 Feb 25. PubMed PMID: 26914441. <https://www.ncbi.nlm.nih.gov/pubmed/26914441>

Ma Z, Moruzzi N, Catrina SB, Grill V, Björklund A. *Hyperoxia inhibits glucose-induced insulin secretion and mitochondrial metabolism in rat pancreatic islets*. Biochem Biophys Res Commun. 2014 Jan 3;443(1): 223-8. doi: 10.1016/j.bbrc.2013.11.088. Epub 2013 Dec 2. PubMed PMID: 24299957. <https://www.ncbi.nlm.nih.gov/pubmed/24299957>

Gao CL, Liu GL, Liu S, Chen XH, Ji CB, Zhang CM, Xia ZK, Guo XR. *Overexpression of PGC-1 β improves insulin sensitivity and mitochondrial function in 3T3-L1 adipocytes.* Mol Cell Biochem. 2011 Jul;353(1-2):215-23. doi: 10.1007/s11010-011-0789-2. Epub 2011 Apr 16. PubMed PMID: 21499715. <https://www.ncbi.nlm.nih.gov/pubmed/21499715>

Brown-Borg HM, Johnson WT, Rakoczy SG. *Expression of oxidative phosphorylation components in mitochondria of long-living Ames dwarf mice.* Age (Dordr). 2012 Feb;34(1):43-57. doi: 10.1007/s11357-011-9212-x. Epub 2011 Feb 16. PubMed PMID: 21327718; PubMed Central PMCID: PMC3260352. <https://www.ncbi.nlm.nih.gov/pubmed/21327718>

Yechoor VK, Patti ME, Saccone R, Kahn CR. *Coordinated patterns of gene expression for substrate and energy metabolism in skeletal muscle of diabetic mice.* Proc Natl Acad Sci U S A. 2002 Aug 6;99(16):10587-92. Epub 2002 Jul 29. PubMed PMID: 12149437; PubMed Central PMCID: PMC124982. <https://www.ncbi.nlm.nih.gov/pubmed/12149437>

Mingrone G, Manco M, Calvani M, Castagneto M, Naon D, Zorzano A. *Could the low level of expression of the gene encoding skeletal muscle mitofusin-2 account for the metabolic inflexibility of obesity?* Diabetologia. 2005 Oct; 48(10): 2108-14. Epub 2005 Sep 14. PubMed PMID: 16160866. DOI: 10.1007/s00125-005-1918-9
<https://www.ncbi.nlm.nih.gov/pubmed/16160866>

Yoshioka S, Okimura Y, Takahashi Y, Iida K, Kaji H, Matsuo M, Chihara K. *Up-regulation of mitochondrial transcription factor 1 mRNA levels by GH in VSMC.* Life Sci. 2004 Mar 12;74(17):2097-109. PubMed PMID: 14969715. DOI: 10.1016/j.lfs.2003.07.057.
<https://www.ncbi.nlm.nih.gov/pubmed/14969715>

3) Complex III Subunit 4:

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+COMPLEX+III+SUBUNIT+4>

Sort by: Most Recent - Search results = Item: 18

Shelley P, Martin-Gronert MS, Rowleron A, Poston L, Heales SJ, Hargreaves IP, McConnell JM, Ozanne SE, Fernandez-Twinn DS. *Altered skeletal muscle insulin signaling and mitochondrial complex II-III linked activity in adult offspring of obese mice.* Am J Physiol Regul Integr Comp Physiol. 2009 Sep; 297(3) :R675-81. doi: 10.1152/ajpregu.00146.2009. Epub 2009 Jun 17. PubMed PMID: 19535678; PubMed Central PMCID: PMC2739782.
<https://www.ncbi.nlm.nih.gov/pubmed/19535678>

Lee B, Srinivasan M, Aalinkeel R, Patel MS, Laychock SG. *Mitochondrial-encoded gene regulation in rat pancreatic islets.* Lee Metabolism. 2001 Feb;50(2):200-6. PubMed PMID: 11229430. DOI: 10.1053/meta.2001.17714. <https://www.ncbi.nlm.nih.gov/pubmed/11229430>

4) Complex III Subunit 4:

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+COMPLEX+III+SUBUNIT+4>

Sort by: Best Match - Search results = Item: 12

Balhara B, Burkart A, Topcu V, Lee YK, Cowan C, Kahn CR, Patti ME. *Severe insulin resistance alters metabolism in mesenchymal progenitor cells.* *Endocrinology.* 2015 Jun;156(6):2039-48. doi: 10.1210/en.2014-1403. Epub 2015 Mar 26. PubMed PMID: 25811318; PubMed Central PMCID: PMC4430624. <https://www.ncbi.nlm.nih.gov/pubmed/25811318>

Heilbronn LK, Gan SK, Turner N, Campbell LV, Chisholm DJ. *Markers of mitochondrial biogenesis and metabolism are lower in overweight and obese insulin-resistant subjects.* *J Clin Endocrinol Metab.* 2007 Apr;92(4):1467-73. Epub 2007 Jan 23. PubMed PMID: 17244782. <https://www.ncbi.nlm.nih.gov/pubmed/17244782>

1.31 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS E CYTOCHROME C (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “*Pseudomonas*” e “*Cytochrome c*”.

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+CYTOCHROME+C>

Sort by: Most Recent - Search results = Items: 726

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi**:

Jo J, Cortez KL, Cornell WC, Price-Whelan A, Dietrich LE. *An orphan cbb(3)-type cytochrome oxidase subunit supports Pseudomonas aeruginosa biofilm growth and virulence.* Elife. 2017 Nov 21;6. pii: e30205. doi: 10.7554/eLife.30205. PubMed PMID: 29160206.

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Calero P, Jensen SI, Bojanovič K, Lennen RM, Koza A, Nielsen AT. *Genome-wide identification of tolerance mechanisms toward p-coumaric acid in Pseudomonas putida.* Biotechnol Bioeng. 2017 Nov 13. doi: 10.1002/bit.26495. [Epub ahead of print] PubMed PMID: 29131301.

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Albada B, Metzler-Nolte N. *Highly Potent Antibacterial Organometallic Peptide Conjugates.* Acc Chem Res. 2017 Oct 17; 50(10): 2510-2518. doi: 10.1021/acs.accounts.7b00282. Epub 2017 Sep 27. PubMed PMID: 28953347. <https://www.ncbi.nlm.nih.gov/pubmed/28953347>

Jiang T, Guo X, Yan J, Zhang Y, Wang Y, Zhang M, Sheng B, Ma C, Xu P, Gao C. *A Bacterial Multidomain NAD-Independent d-Lactate Dehydrogenase Utilizes Flavin Adenine Dinucleotide and Fe-S Clusters as Cofactors and Quinone as an Electron Acceptor for d-Lactate Oxidization.* J Bacteriol. 2017 Oct 17;199(22). pii: e00342-17. doi: 10.1128/JB.00342-17. Print 2017 Nov 15. PubMed PMID: 28847921; PubMed Central PMCID: PMC5648861.

<https://www.ncbi.nlm.nih.gov/pubmed/28847921>

Pinck S, Etienne M, Dossot M, Jorand FPA. *A rapid and simple protocol to prepare a living biocomposite that mimics electroactive biofilms.* Bioelectrochemistry. 2017 Dec; 118: 131-138. doi: 10.1016/j.bioelechem.2017.07.010. Epub 2017 Aug 1. PubMed PMID: 28802176.

<https://www.ncbi.nlm.nih.gov/pubmed/28802176>

Osamura T, Kawakami T, Kido R, Ishii M, Arai H. *Specific expression and function of the A-type cytochrome c oxidase under starvation conditions in Pseudomonas aeruginosa.* PLoS One. 2017 May 18;12(5):e0177957. doi: 10.1371/journal.pone.0177957. eCollection 2017. PubMed PMID: 28542449; PubMed Central PMCID: PMC5436846.

<https://www.ncbi.nlm.nih.gov/pubmed/28542449>

Baldissera MD, Souza CF, Grings M, Parmeggiani BS, Leipnitz G, Moreira KLS, da Rocha MIUM, da Veiga ML, Santos RCV, Stefani LM, Baldisserotto B. *Inhibition of the mitochondrial respiratory chain in gills of Rhamdia quelen experimentally infected by Pseudomonas aeruginosa: Interplay with reactive oxygen species.* Microb Pathog. 2017 Jun;107:349-353. doi: 10.1016/j.micpath.2017.04.017. Epub 2017 Apr 13. PubMed PMID: 28414167. <https://www.ncbi.nlm.nih.gov/pubmed/28414167>

Carvalheda CA, Pisljakov AV. *Insights into proton translocation in cbb(3) oxidase from MD simulations.* Biochim Biophys Acta. 2017 May; 1858(5): 396-406. doi: 10.1016/j.bbabi.2017.02.013. Epub 2017 Mar 1. PubMed PMID: 28259641. <https://www.ncbi.nlm.nih.gov/pubmed/28259641>

Kohlstaedt M, Buschmann S, Langer JD, Xie H, Michel H. *Subunit CcoQ is involved in the assembly of the Cbb(3)-type cytochrome c oxidases from pseudomonas stutzeri ZoBell but not required for their activity.* Biochim Biophys Acta. 2017 Mar;1858(3):231-238. doi: 10.1016/j.bbabi.2016.12.006. Epub 2016 Dec 20. PubMed PMID: 28007379. <https://www.ncbi.nlm.nih.gov/pubmed/28007379>

Hirai T, Osamura T, Ishii M, Arai H. *Expression of multiple cbb3 cytochrome c oxidase isoforms by combinations of multiple isosubunits in Pseudomonas aeruginosa.* Proc Natl Acad Sci U S A. 2016 Oct 24. pii: 201613308. [Epub ahead of print] PubMed PMID: 27791152; PubMed Central PMCID: PMC5111723. <https://www.ncbi.nlm.nih.gov/pubmed/27791152>

Melin F, Xie H, Meyer T, Ahn YO, Gennis RB, Michel H, Hellwig P. *The unusual redox properties of C-type oxidases.* Biochim Biophys Acta. 2016 Dec;1857(12):1892-1899. doi: 10.1016/j.bbabi.2016.09.009. Epub 2016 Sep 21. PubMed PMID: 27664317. <https://www.ncbi.nlm.nih.gov/pubmed/27664317>

Kohlstaedt M, Buschmann S, Xie H, Resemann A, Warkentin E, Langer JD, Michel H. *Identification and Characterization of the Novel Subunit CcoM in the cbb3₃ Cytochrome c Oxidase from Pseudomonas stutzeri ZoBell.* MBio. 2016 Jan 26;7(1): e01921-15. doi: 10.1128/mBio.01921-15. PubMed PMID: 26814183; PubMed Central PMCID: PMC4742706. <https://www.ncbi.nlm.nih.gov/pubmed/26814183>

Hazan R, Que YA, Maura D, Strobel B, Majcherczyk PA, Hopper LR, Wilbur DJ, Hreha TN, Barquera B, Rahme LG. *Auto Poisoning of the Respiratory Chain by a Quorum-Sensing-Regulated Molecule Favors Biofilm Formation and Antibiotic Tolerance.* Curr Biol. 2016 Jan 25;26(2):195-206. doi: 10.1016/j.cub.2015.11.056. Epub 2016 Jan 14. PubMed PMID: 26776731; PubMed Central PMCID: PMC4729643. <https://www.ncbi.nlm.nih.gov/pubmed/26776731>

Wood SJ, Goldufsky JW, Bello D, Masood S, Shafikhani SH. *Pseudomonas aeruginosa ExoT Induces Mitochondrial Apoptosis in Target Host Cells in a Manner That Depends on Its GTPase-activating Protein (GAP) Domain Activity.* J Biol Chem. 2015 Nov 27;290(48):29063-73. doi: 10.1074/jbc.M115.689950. Epub 2015 Oct 8. PubMed PMID: 26451042; PubMed Central PMCID: PMC4661418. <https://www.ncbi.nlm.nih.gov/pubmed/26451042>

Veena VK, Popavath RN, Kennedy K, Sakthivel N. *In vitro antiproliferative, pro-apoptotic, antimetastatic and anti-inflammatory potential of 2,4-diacetylphloroglucinol (DAPG) by Pseudomonas aeruginosa strain FP10.* Apoptosis. 2015 Oct;20(10):1281-95. doi: 10.1007/s10495-015-1162-9. PubMed PMID: 26283170. <https://www.ncbi.nlm.nih.gov/pubmed/26283170>

Matsuno T, Yumoto I. *Bioenergetics and the role of soluble cytochromes C for alkaline adaptation in gram-negative alkaliphilic Pseudomonas.* Biomed Res Int. 2015;2015:847945. doi: 10.1155/2015/847945. Epub 2015 Feb 2. Review. PubMed PMID: 25705691; PubMed Central PMCID: PMC4332470. <https://www.ncbi.nlm.nih.gov/pubmed/25705691>

Managò A, Becker KA, Carpinteiro A, Wilker B, Soddemann M, Seitz AP, Edwards MJ, Grassmé H, Szabò I, Gulbins E. *Pseudomonas aeruginosa pyocyanin induces neutrophil death via mitochondrial reactive oxygen species and mitochondrial acid sphingomyelinase.* Antioxid Redox Signal. 2015 May 1;22(13):1097-110. doi: 10.1089/ars.2014.5979. Epub 2015 Mar 18. PubMed PMID: 25686490; PubMed Central PMCID: PMC4403017. <https://www.ncbi.nlm.nih.gov/pubmed/25686490>

Levin BD, Walsh KA, Sullivan KK, Bren KL, Elliott SJ. *Methionine ligand lability of homologous monoheme cytochromes c.* Inorg Chem. 2015 Jan 5;54(1):38-46. doi: 10.1021/ic501186h. Epub 2014 Dec 9. PubMed PMID: 25490149. <https://www.ncbi.nlm.nih.gov/pubmed/25490149>

Arai H, Kawakami T, Osamura T, Hirai T, Sakai Y, Ishii M. *Enzymatic characterization and in vivo function of five terminal oxidases in Pseudomonas aeruginosa.* J Bacteriol. 2014 Dec;196(24):4206-15. doi: 10.1128/JB.02176-14. Epub 2014 Sep 2. PubMed PMID: 25182500; PubMed Central PMCID: PMC4248849. <https://www.ncbi.nlm.nih.gov/pubmed/25182500>

Hamada M, Toyofuku M, Miyano T, Nomura N. *cbb3-type cytochrome c oxidases, aerobic respiratory enzymes, impact the anaerobic life of Pseudomonas aeruginosa PAO1.* J Bacteriol. 2014 Nov;196(22):3881-9. doi: 10.1128/JB.01978-14. Epub 2014 Sep 2. PubMed PMID: 25182494; PubMed Central PMCID: PMC4248832. <https://www.ncbi.nlm.nih.gov/pubmed/25182494>

1.32 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS E CYTOCHROME C (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle migliori corrispondenze tra i termini “*Pseudomonas*” e “*Cytochrome c*” (**Attenzione:** gli articoli già citati nella ricerca precedente non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+CYTOCHROME+C>

Sort by: Best Match - Search results = Item: 634

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Schwarzer C, Fu Z, Shuai S, Babbar S, Zhao G, Li C, Machen TE. *Pseudomonas aeruginosa* homoserine lactone triggers apoptosis and Bak/Bax-independent release of mitochondrial cytochrome *C* in fibroblasts. *Cell Microbiol.* 2014 Jul;16(7):1094-104. doi: 10.1111/cmi.12263. Epub 2014 Feb 13. PubMed PMID: 24438098; PubMed Central PMCID: PMC4065231. <https://www.ncbi.nlm.nih.gov/pubmed/24438098>

Sun Y, Benabbas A, Zeng W, Kleingardner JG, Bren KL, Champion PM. *Investigations of heme distortion, low-frequency vibrational excitations, and electron transfer in cytochrome c.* *Proc Natl Acad Sci U S A.* 2014 May 6;111(18):6570-5. doi: 10.1073/pnas.1322274111. Epub 2014 Apr 21. PubMed PMID: 24753591; PubMed Central PMCID: PMC4020103. <https://www.ncbi.nlm.nih.gov/pubmed/24753591>

Xie H, Buschmann S, Langer JD, Ludwig B, Michel H. *Biochemical and biophysical characterization of the two isoforms of cbb3-type cytochrome c oxidase from Pseudomonas stutzeri.* *J Bacteriol.* 2014 Jan;196(2):472-82. doi: 10.1128/JB.01072-13. Epub 2013 Nov 8. PubMed PMID: 24214947; PubMed Central PMCID: PMC3911263. <https://www.ncbi.nlm.nih.gov/pubmed/24214947>

Buschmann S, Richers S, Ermler U, Michel H. *A decade of crystallization drops: crystallization of the cbb3 cytochrome c oxidase from Pseudomonas stutzeri.* *Protein Sci.* 2014 Apr;23(4):411-22. doi: 10.1002/pro.2423. Epub 2014 Feb 4. PubMed PMID: 24488923; PubMed Central PMCID: PMC3970892. <https://www.ncbi.nlm.nih.gov/pubmed/24488923>

Di Silvio E, Di Matteo A, Malatesta F, Travaglini-Allocatelli C. *Recognition and binding of apocytochrome c to P. aeruginosa CcmI, a component of cytochrome c maturation machinery.* *Biochim Biophys Acta.* 2013 Aug;1834(8):1554-61. doi: 10.1016/j.bbapap.2013.04.027. Epub 2013 May 3. PubMed PMID: 23648553. <https://www.ncbi.nlm.nih.gov/pubmed/23648553>

Matsuno T, Yoshimune K, Yumoto I. *Physiological function of soluble cytochrome c-552 from alkaliphilic Pseudomonas alcaliphila AL15-21(T).* *J Bioenerg Biomembr.* 2011 Oct;43(5):473-81. doi: 10.1007/s10863-011-9376-1. Epub 2011 Jul 16. PubMed PMID: 21766198. <https://www.ncbi.nlm.nih.gov/pubmed/21766198>

Matsuno T, Mie Y, Yoshimune K, Yumoto I. *Physiological role and redox properties of a small cytochrome c(5), cytochrome c-552, from alkaliphile, Pseudomonas alcaliphila AL15-21(T).* *J Biosci Bioeng.* 2009 Dec;108(6):465-70. doi: 10.1016/j.jbiosc.2009.06.008. Epub 2009 Jul 14. PubMed PMID: 19914577. <https://www.ncbi.nlm.nih.gov/pubmed/19914577>

Liu X, Tremblay PL, Malvankar NS, Nevin KP, Lovley DR, Vargas M. *A Geobacter sulfurreducens strain expressing pseudomonas aeruginosa type IV pili localizes OmcS on pili but is deficient in Fe(III) oxide reduction and current production.* *Appl Environ Microbiol.* 2014 Feb;80(3):1219-24. doi: 10.1128/AEM.02938-13. Epub 2013 Dec 2. PubMed PMID: 24296506; PubMed Central PMCID: PMC3911229. <https://www.ncbi.nlm.nih.gov/pubmed/24296506>

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Needleman SB, Blair TT. *Homology of Pseudomonas cytochrome c-551 with eukaryotic c-cytochromes.* Proc Natl Acad Sci U S A. 1969 Aug;63(4):1227-33. PubMed PMID: 5260924; PubMed Central PMCID: PMC223454. <https://www.ncbi.nlm.nih.gov/pubmed/5260924>

1.33 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS E CYTOCHROME C1 (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Pseudomonas*” e “*Cytochrome c1*”.

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+CYTOCHROME+C1>

Sort by: Most Recent - Search results = Items: 18

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi**:

García I, Rosas T, Bejarano ER, Gotor C, Romero LC. *Transient transcriptional regulation of the CYS-C1 gene and cyanide accumulation upon pathogen infection in the plant immune response.* Plant Physiol. 2013 Aug;162(4):2015-27. doi: 10.1104/pp.113.219436. Epub 2013 Jun 19. PubMed PMID: 23784464; PubMed Central PMCID: PMC3729779.

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Ben-Bassat A, Goldberg I. *Oxidation of C1-compounds in Pseudomonas C.* Biochim Biophys Acta. 1977 Apr 27;497(2):586-97. PubMed PMID: 192317. <https://www.ncbi.nlm.nih.gov/pubmed/192317>

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Wilson MT, Greenwood C, Brunori M, Antonini E. *Electron transfer between azurin and cytochrome c-551 from Pseudomonas aeruginosa.* Biochem J. 1975 Mar;145(3):449-57. PubMed PMID: 168867; PubMed Central PMCID: PMC1165244. <https://www.ncbi.nlm.nih.gov/pubmed/168867>

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1.34 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS E CYTOCHROME C1 (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle migliori corrispondenze tra i termini “*Pseudomonas*” e “*Cytochrome c1*” (**Attenzione:** gli articoli già citati nella ricerca precedente non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+CYTOCHROME+C1>

Sort by: Best Match - Search results = Item: 20

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Yamaguchi M, Fujisawa H. *Reconstitution of iron-sulfur cluster of NADH-cytochrome c reductase, a component of benzoate 1,2-dioxygenase system from Pseudomonas arvilla C-1.* J Biol Chem. 1981 Jul 10;256(13):6783-7. PubMed PMID: 7240244.

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1. 35 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA BURKHOLDERIA E CYTOCHROME C (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “*Burkholderia*” e “*Cytochrome c*”.

<https://www.ncbi.nlm.nih.gov/pubmed/?term=BURKHOLDERIA+CYTOCHROME+C>

Sort by: Most Recent - Search results = Items: 13

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi**:

Jones-Carson J, Zweifel AE, Tapscott T, Austin C, Brown JM, Jones KL, Voskuil MI, Vázquez-Torres A. Nitric oxide from IFN γ -primed macrophages modulates the antimicrobial activity of β -lactams against the intracellular pathogens *Burkholderia pseudomallei* and *Nontyphoidal Salmonella*. PLoS Negl Trop Dis. 2014 Aug 14;8(8):e3079. doi: 10.1371/journal.pntd.0003079. eCollection 2014 Aug. PubMed PMID: 25121731; PubMed Central PMCID: PMC4133387. <https://www.ncbi.nlm.nih.gov/pubmed/25121731>

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1. 36 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA BURKHOLDERIA E CYTOCHROME C1

<https://www.ncbi.nlm.nih.gov/pubmed/?term=BURKHOLDERIA+CYTOCHROME+C1>

Search results = Items: 1

Kang MJ, Lee MH, Shim JK, Seo ST, Shrestha R, Cho MS, Hahn JH, Park DS. *PCR-based specific detection of Ralstonia solanacearum by amplification of cytochrome c1 signal peptide sequences.* *J Microbiol Biotechnol.* 2007 Nov;17(11):1765-71. PubMed PMID: 18092459.

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1.37 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E TRANSPORTER, (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Transporter*”.

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+TRANSPORTER>

Sort by: Most Recent - Search results = Items: 76

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Stechemesser L, Eder SK, Wagner A, Patsch W, Feldman A, Strasser M, Auer S, Niederseer D, Huber-Schönauer U, Paulweber B, Zandanell S, Ruhaltinger S, Weghuber D, Haschke-Becher E, Grabmer C, Rohde E, Datz C, Felder TK, Aigner E. *Metabolomic profiling identifies potential pathways involved in the interaction of iron homeostasis with glucose metabolism.* Mol Metab. 2016 Oct 31;6(1):38-47. doi: 10.1016/j.molmet.2016.10.006. eCollection 2017 Jan. PubMed PMID: 28123936; PubMed Central PMCID: PMC5220278.

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<https://www.ncbi.nlm.nih.gov/pubmed/22450556>

Pollak Y, Mechlovich D, Amit T, Bar-Am O, Manov I, Mandel SA, Weinreb O, Meyron-Holtz EG, Iancu TC, Youdim MB. *Effects of novel neuroprotective and neurorestorative multifunctional drugs on iron chelation and glucose metabolism.* *J Neural Transm (Vienna).* 2013 Jan;120(1):37-48. doi: 10.1007/s00702-012-0795-x. Epub 2012 Mar 25. PubMed PMID: 22446839.
<https://www.ncbi.nlm.nih.gov/pubmed/22446839>

1.38 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E TRANSPORTER, (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *migliori* corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Transporter*” (**Attenzione:** gli articoli già citati nella ricerca precedente non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+TRANSPORTER>

Sort by: Best Match - Search results = Item: 68

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi**:

Dongiovanni P, Valenti L, Ludovica Fracanzani A, Gatti S, Cairo G, Fargion S. *Iron depletion by deferoxamine up-regulates glucose uptake and insulin signaling in hepatoma cells and in rat liver.* Am J Pathol. 2008 Mar;172(3):738-47. doi: 10.2353/ajpath.2008.070097. Epub 2008 Feb 2. PubMed PMID: 18245813; PubMed Central PMCID: PMC2258266.
<https://www.ncbi.nlm.nih.gov/pubmed/18245813>

Datz C, Felder TK, Niederseer D, Aigner E. *Iron homeostasis in the metabolic syndrome.* Eur J Clin Invest. 2013 Feb;43(2):215-24. doi: 10.1111/eci.12032. Epub 2013 Jan 7. Review. PubMed PMID: 23289518. <https://www.ncbi.nlm.nih.gov/pubmed/23289518>

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1.39 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E DIABETE (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Transporter*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+DIABETES>

Sort by: Most Recent - Search results = Items: 650

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

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1.40 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E DIABETE (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *migliori* corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Diabete*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+DIABETES>

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Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Sen S, Ghatak SK, Majumdar D, Sen K, Bhattacharya B. *Free iron status & insulin resistance in type 2 diabetes mellitus: Analyzing the probable role of a peanut protein.* Indian J Med Res. 2015 Nov;142(5):606-9. doi: 10.4103/0971-5916.171291. PubMed PMID: 26658597; PubMed Central PMCID: PMC4743349. <https://www.ncbi.nlm.nih.gov/pubmed/26658597>

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1.41 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E OBESITA' (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Obesità*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+OBESITY>

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Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

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1.42 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E OBESITA' (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *migliori* corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Obesità*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+OBESITY>

Sort by: Best Match - Search results = Items: 189

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

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1.43 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E INSULITE (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Insulite*”.

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+INSULITIS>

Sort by: Most Recent - Search results = Items: 4

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1.44 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E FIBROSI CISTICA (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Fibrosi Cistica*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+CYSTIC+FIBROSI>
[S](#)

Sort by: Most Recent - Search results = Items: 4

Kessler L, Abély M. [*Pancreatic infringement exocrine and endocrine in cystic fibrosis*]. Arch Pediatr. 2016 Dec;23(12S):12S21-12S32. doi: 10.1016/S0929-693X(17)30059-3. French. PubMed PMID: 28231890. <https://www.ncbi.nlm.nih.gov/pubmed/28231890>

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1.45 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E INFEZIONI POLMONARI (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Infezioni Polmonari*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

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Sort by: Most Recent - Search results = Items: 4

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1.46 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E SISTEMA IMMUNITARIO (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Sistema Immunitario*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+IMMUNE+SYSTEM>

Sort by: Most Recent - Search results = Items: 57

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

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1.47 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E APOPTOSI (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Apoptosi*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+APOPTOSIS>

Sort by: Most Recent - Search results = Items: 57

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Backe MB, Moen IW, Ellervik C, Hansen JB, Mandrup-Poulsen T. *Iron Regulation of Pancreatic Beta-Cell Functions and Oxidative Stress.* *Annu Rev Nutr.* 2016 Jul 17;36:241-73. doi: 10.1146/annurev-nutr-071715-050939. Epub 2016 May 4. Review. PubMed PMID: 27146016.
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1.48 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E APOPTOSI (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *migliori* corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Apoptosi*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+APOPTOSIS>

Sort by: Best Match - Search results = Items: 48

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

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1.49 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E CYTOCHROME C (UNA SELEZIONE DELLE *PIU' RECENTI* CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Cytochrome c*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+CYTOCHROME+C>

Sort by: Most Recent - Search results = Items: 16

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Zou C, Liu X, Xie R, Bao Y, Jin Q, Jia X, Li L, Liu R. *Deferiprone attenuates inflammation and myocardial fibrosis in diabetic cardiomyopathy rats.* Biochem Biophys Res Commun. 2017 May 13;486(4):930-936. doi: 10.1016/j.bbrc.2017.03.127. Epub 2017 Mar 24. PubMed PMID: 28347819. <https://www.ncbi.nlm.nih.gov/pubmed/28347819>

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1.50 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E LE REACTIVE OXYGEN SPECIES (UNA SELEZIONE DELLE *PIU' RECENTI* CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Reactive Oxygen Species*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+REACTIVE+OXYGEN+SPECIES>

Sort by: Most Recent - Search results = Items: 92

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

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1.51 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA, IRON E LE REACTIVE OXYGEN SPECIES (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *migliori* corrispondenze tra i termini “*Insulina*”, “*Iron*” e “*Reactive Oxygen Species*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+IRON+REACTIVE+OXYGEN+SPECIES>

Sort by: Best Match - Search results = Items: 78

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

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1.52 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E TRANSPORTER (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Transporter*”.

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+IRON+TRANSPORTER>

Sort by: Most Recent - Search results = Items: 247

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1.53 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E TRANSPORTER (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *migliori* corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Transporter*” (**Attenzione:** gli articoli già citati nella ricerca precedente non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+IRON+TRANSPORTER>

Sort by: Best Match - Search results = Items: 212

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

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1.54 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E MELIOIDOSI (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “Pseudomonas”, “Iron” e “Meliodosi” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+IRON+MELIOIDOSIS>

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Wanachiwanawin W. *Infections in E-beta thalassemia.* J Pediatr Hematol Oncol. 2000 Nov-Dec;22(6):581-7. Review. PubMed PMID: 11132234.
<https://www.ncbi.nlm.nih.gov/pubmed/11132234>

Sexton MM, Jones AL, Chaowagul W, Woods DE. *Purification and characterization of a protease from Pseudomonas pseudomallei.* Can J Microbiol. 1994 Nov;40(11):903-10. PubMed PMID: 7528633. <https://www.ncbi.nlm.nih.gov/pubmed/7528633>

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1.55 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E DIABETE (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Diabete*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

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Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Lakatos B, Nikolova R, Ocskay L, Csomor J, Prinz G. [Case of a diabetic man cured of rhinocerebral zygomycosis]. *Orv Hetil.* 2010 Sep 26;151(39):1591-6. doi: 10.1556/OH.2010.28969. Hungarian. PubMed PMID: 20840916.
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1.56 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E FIBROSI CISTICA (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Fibrosi Cistica*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+IRON+CYSTIC+FIBROSIS>

Sort by: Most Recent - Search results = Items: 148

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi**:

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Kang D, Kirienko NV. *High-Throughput Genetic Screen Reveals that Early Attachment and Biofilm Formation Are Necessary for Full Pyoverdine Production by Pseudomonas aeruginosa.* Front Microbiol. 2017 Sep 5;8:1707. doi: 10.3389/fmicb.2017.01707. eCollection 2017. PubMed PMID: 28928729; PubMed Central PMCID: PMC5591869. <https://www.ncbi.nlm.nih.gov/pubmed/28928729>

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1.57 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E FIBROSI CISTICA (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *migliori* corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Fibrosi Cistica*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+IRON+CYSTIC+FIBROSIS>

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Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Nguyen AT, O'Neill MJ, Watts AM, Robson CL, Lamont IL, Wilks A, Oglesby-Sherrouse AG. *Adaptation of iron homeostasis pathways by a Pseudomonas aeruginosa pyoverdine mutant in the cystic fibrosis lung.* J Bacteriol. 2014 Jun;196(12):2265-76. doi: 10.1128/JB.01491-14. Epub 2014 Apr 11. PubMed PMID: 24727222; PubMed Central PMCID: PMC4054187. <https://www.ncbi.nlm.nih.gov/pubmed/24727222>

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1.58 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E INFEZIONI POLMONARI (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Infezioni Polmonari*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+IRON+PULMONARY+INFECTION>

Sort by: Most Recent - Search results = Items: 74

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi**:

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1.59 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E INFEZIONI POLMONARI (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *migliori* corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Infezioni Polmonari*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+IRON+PULMONARY+INFECTION>

Sort by: Best Match - Search results = Items: 100

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

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1.60 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E SISTEMA IMMUNITARIO (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Sistema Immunitario*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+IRON+IMMUNE+SYSTEM>

Sort by: Most Recent - Search results = Items: 63

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

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1.61 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E SISTEMA IMMUNITARIO (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *migliori* corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Sistema Immunitario*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

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Sort by: Best Match - Search results = Items: 62

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi**:

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1.62 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E APOPTOSI (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Apoptosi*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+IRON+APOPTOSIS>

Sort by: Most Recent - Search results = Items: 8

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Lim CJ, Kim WB, Lee BS, Lee HY, Kwon TH, Park JM, Kwon SY. *Silencing of SIFTR-c, the catalytic subunit of ferredoxin:thioredoxin reductase, induces pathogenesis-related genes and pathogen resistance in tomato plants.* Biochem Biophys Res Commun. 2010 Sep 3;399(4):750-4. doi: 10.1016/j.bbrc.2010.08.016. Epub 2010 Aug 10. PubMed PMID: 20705057.

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1.63 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E LE REACTIVE OXYGEN SPECIES (UNA SELEZIONE DELLE *PIU' RECENTI* CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Reactive Oxygen Species*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+IRON+REACTIVE+OXYGEN+SPECIES>

Sort by: Most Recent - Search results = Items: 111

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Pasqua M, Visaggio D, Lo Sciuto A, Genah S, Banin E, Visca P, Imperi F. *Ferric Uptake Regulator Fur Is Conditionally Essential in Pseudomonas aeruginosa.* J Bacteriol. 2017 Oct 17;199(22). pii: e00472-17. doi: 10.1128/JB.00472-17. Print 2017 Nov 15. PubMed PMID: 28847923; PubMed Central PMCID: PMC5648857.

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1.64 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E LE REACTIVE OXYGEN SPECIES (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *migliori* corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Reactive Oxygen Species*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+IRON+REACTIVE+OXYGEN+SPECIES>

Sort by: Best Match - Search results = Items: 108

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Rodríguez-Chueca J, Morales M, Mosteo R, Ormad MP, Ovelleiro JL. *Inactivation of Enterococcus faecalis, Pseudomonas aeruginosa and Escherichia coli present in treated urban wastewater by coagulation-flocculation and photo-Fenton processes.* Photochem Photobiol Sci. 2013 May;12(5):864-71. doi: 10.1039/c3pp25352j. PubMed PMID: 23411627. <https://www.ncbi.nlm.nih.gov/pubmed/23411627>

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Yamakura F, Suzuki K. *Inactivation of Pseudomonas iron-superoxide dismutase by hydrogen peroxide.* Biochim Biophys Acta. 1986 Nov 7;874(1):23-9. PubMed PMID: 3768375. <https://www.ncbi.nlm.nih.gov/pubmed/3768375>

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Ahmad SI, Iranzo OG. *Treatment of post-burns bacterial infections by Fenton reagent, particularly the ubiquitous multiple drug resistant Pseudomonas spp.* Med Hypotheses. 2003 Oct;61(4):431-4. PubMed PMID: 13679006. <https://www.ncbi.nlm.nih.gov/pubmed/13679006>

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Avellan A, Auffan M, Masion A, Levard C, Bertrand M, Rose J, Santaella C, Achouak W. *Remote Biodegradation of Ge-Imogolite Nanotubes Controlled by the Iron Homeostasis of Pseudomonas brassicacearum.* Environ Sci Technol. 2016 Jul 19;50(14):7791-8. doi: 10.1021/acs.est.6b01455. Epub 2016 Jun 27. PubMed PMID: 27347687.

<https://www.ncbi.nlm.nih.gov/pubmed/27347687>

Park SC, Kim NH, Yang W, Nah JW, Jang MK, Lee D. *Polymeric micellar nanoplatfoms for Fenton reaction as a new class of antibacterial agents.* J Control Release. 2016 Jan 10;221:37-47. doi: 10.1016/j.jconrel.2015.11.027. Epub 2015 Dec 2. PubMed PMID: 26639177.

<https://www.ncbi.nlm.nih.gov/pubmed/26639177>

1.65 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E CYTOCHROME C (UNA SELEZIONE DELLE *PIU' RECENTI* CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Cytochrome c*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=PSEUDOMONAS+IRON+CYTOCHROME+C>

Sort by: Most Recent - Search results = Items: 108

Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

Jiang T, Guo X, Yan J, Zhang Y, Wang Y, Zhang M, Sheng B, Ma C, Xu P, Gao C. *A Bacterial Multidomain NAD-Independent d-Lactate Dehydrogenase Utilizes Flavin Adenine Dinucleotide and Fe-S Clusters as Cofactors and Quinone as an Electron Acceptor for d-Lactate Oxidization.* J Bacteriol. 2017 Oct 17;199(22). pii: e00342-17. doi: 10.1128/JB.00342-17. Print 2017 Nov 15. PubMed PMID: 28847921; PubMed Central PMCID: PMC5648861.

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1.66 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA PSEUDOMONAS, IRON E CYTOCHROME C (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle migliori corrispondenze tra i termini “*Pseudomonas*”, “*Iron*” e “*Cytochrome c*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

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Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

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1.67 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA BURKHOLDERIA, IRON E TRANSPORTER (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “*Burkholderia*”, “*Iron*” e “*Transporter*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=BURKHOLDERIA+IRON+TRANSPORTER>

Sort by: Most Recent - Search results = Items: 25

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1.68 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA BURKHOLDERIA, IRON E TRANSPORTER (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle migliori corrispondenze tra i termini “*Burkholderia*”, “*Iron*” e “*Transporter*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=BURKHOLDERIA+IRON+TRANSPORTER>

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Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi:**

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1.69 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA BURKHOLDERIA, IRON E FIBROSI CISTICA (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

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Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi**:

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1.70 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA BURKHOLDERIA, IRON E SISTEMA IMMUNITARIO (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Burkholderia*”, “*Iron*” e “*Immune System*”.

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Sort by: Most Recent - Search results = Items: 4

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1.71 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA BURKHOLDERIA, IRON E CYTOCHROME C (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Burkholderia*”, “*Iron*” e “*Cytochrome c*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

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1.72 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA BURKHOLDERIA, IRON E LE REACTIVE OXYGEN SPECIES (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Burkholderia*”, “*Iron*” e “*Reactive Oxygen Species*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

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1.73 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA BURKHOLDERIA, IRON E MELIOIDOSI (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

I **Link** mostrati qui di seguito effettuano ricerche bibliografiche (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Burkholderia*”, “*Iron*” e “*Melioidosi*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

<https://www.ncbi.nlm.nih.gov/pubmed/?term=BURKHOLDERIA+IRON+MELIOIDOSIS>

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Dai risultati della ricerca sono stati selezionati **alcuni articoli ritenuti significativi**:

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BIBLIOGRAFIA RIGUARDANTE HELIGMOSOMOIDES POLYGYRUS

1. 74 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA INSULINA E HELIGMOSOMOIDES POLYGYRUS (UNA SELEZIONE DELLE *PIU' RECENTI* CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Insulin*” e “*Heligmosomoides polygyrus*”.

<https://www.ncbi.nlm.nih.gov/pubmed/?term=INSULIN+HELIGMOSOMOIDES+POLYGYRUS>

Sort by: Most Recent - Search results = Items: 5

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1.75 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA HELIGMOSOMOIDES POLYGYRUS E SISTEMA IMMUNITARIO (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle più recenti corrispondenze tra i termini “*Heligmosomoides polygyrus*” e “*Immune system*”.

<https://www.ncbi.nlm.nih.gov/pubmed/?term=HELIGMOSOMOIDES+POLYGYRUS+IMMUNE+SYSTEM>

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Considerato il notevole numero di articoli trovati, allo scopo di restringere maggiormente il campo di ricerca, dai risultati ottenuti sono stati selezionati gli **articoli ritenuti più significativi** attraverso un'ulteriore inserimento di **termini specifici**. (**Attenzione: gli articoli già citati nella ricerca precedente non vengono citati**).

1) “*Heligmosomoides polygyrus, Immune system, TH2, CD4, IL-4*”:

<https://www.ncbi.nlm.nih.gov/pubmed/?term=HELIGMOSOMOIDES+POLYGYRUS+IMMUNE+SYSTEM+TH2+CD4+IL+4>

Sort by: Most Recent - Search results = Items: 26

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1.76 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA HELIGMOSOMOIDES POLYGYRUS E APOPTOSI (UNA SELEZIONE DELLE *PIU' RECENTI* CORRISPONDENZE)

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1.77 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA HELIGMOSOMOIDES POLYGYRUS E INFEZIONI POLMONARI (UNA SELEZIONE DELLE *PIU' RECENTI* CORRISPONDENZE)

Il **Link** mostrato qui di seguito effettua una ricerca bibliografica (su tutti gli **articoli scientifici** pubblicati su **PubMed**) delle *più recenti* corrispondenze tra i termini “*Heligmosomoides polygyrus*” e “*Pulmonary Infection*” (**Attenzione:** gli articoli già citati nelle ricerche precedenti non vengono citati).

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1.78 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA HELIGMOSOMOIDES POLYGYRUS E INFEZIONI POLMONARI (UNA SELEZIONE DELLE MIGLIORI CORRISPONDENZE)

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1.79 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA HELIGMOSOMOIDES POLYGYRUS E DIABETE (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

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1.80 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA HELIGMOSOMOIDES POLYGYRUS E OBESITA' (UNA SELEZIONE DELLE PIU' RECENTI CORRISPONDENZE)

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1.81 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA HELIGMOSOMOIDES POLYGYRUS E CYTOCHROME C (UNA SELEZIONE DELLE *PIU' RECENTI* CORRISPONDENZE)

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<https://www.ncbi.nlm.nih.gov/pubmed/?term=HELIGMOSOMOIDES+POLYGYRUS+CYTOCHROME+C>

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1.82 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA HELIGMOSOMOIDES POLYGYRUS E IRON

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<https://www.ncbi.nlm.nih.gov/pubmed/?term=HELIGMOSOMOIDES+POLYGYRUS+IRON>

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1.83 ARTICOLI SCIENTIFICI CHE STUDIANO LE RELAZIONI TRA HELIGMOSOMOIDES POLYGYRUS E PSEUDOMONAS

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