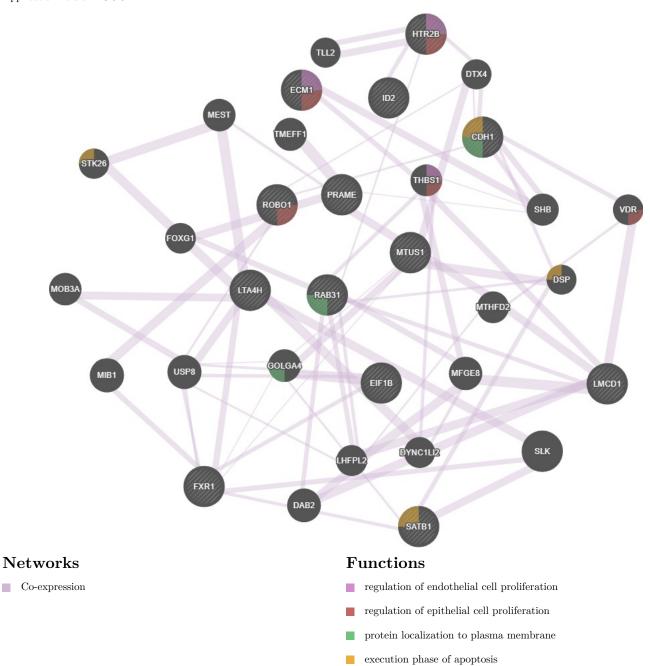
GeneMANIA report

Created on: 12 August 2018 18:05:48

Last database update : 13 March 2017 00:00:00

Application version: 3.6.0



Search parameters

Organism Homo sapiens (human)

Genes CDH1, LMCD1, LTA4H, FXR1, EIF1B, ID2, MTUS1, HTR2B, ECM1,

RAB31, ROBO1, SATB1, PRAME

Network Automatically selected weighting method

weighting

Networks

 \mathbf{A}

Abu-Odeh-Aqeilan-2014 , Agrawal-Sedivy-2010 , Aichem-Groettrup-2012 , Albers-Koegl-2005 , Alexandru-Deshaies-2008 , Alizadeh-Staudt-2000 , Andresen-Flores-Morales-2014 , Arbuckle-Grant-2010 , Arroyo-Aloy-2014 , Arroyo-Aloy-2015

 \mathbf{B}

Bahr-Bowler-2013 , Bailey-Hieter-2015 , Bandyopadhyay-Ideker-2010 , Bantscheff-Drewes-2011 , Barr-Knapp-2009 , Barrios-Rodiles-Wrana-2005 , Behrends-Harper-2010 , Behzadnia-Lührmann-2007 , Bennett-Harper-2010 , Benzinger-Hermeking-2005 , Berggård-James-2006 , Bett-Hay-2013 , Bhatnagar-Attie-2014 , Bild-Nevins-2006 B , BIOGRID-SMALL-SCALE-STUDIES , BIOGRID-SMALL-SCALE-STUDIES , Blandin-Richard-2013 , Blomen-Brummelkamp-2015 , Blomen-Brummelkamp-2015 , Bogachek-Weigel-2014 , Boldrick-Relman-2002 , Bonacci-Soubeyran-2014 , Bouwmeester-Superti-Furga-2004 , Brajenovic-Drewes-2004 , Brehme-Superti-Furga-2009 , Bruderer-Hay-2011 , Burington-Shaughnessy-2008 , Butland-Hayden-2014 , Byron-Humphries-2012

 \mathbf{C}

Cai-Conaway-2007 , Camargo-Brandon-2007 , Campos-Reinberg-2015 , Cao-Chinnaiyan-2014 , Carmon-Liu-2014 , CELL_MAP , Chen-Brown-2002 , Chen-Ge-2013 , Chen-Huang-2014 , Chen-Zhang-2013 , Christianson-Kopito-2011 , Cloutier-Coulombe-2013 , Colland-Gauthier-2004 , Corominas-Iakoucheva-2014 , Couzens-Gingras-2013 , Cox-Rizzino-2013 , Coyaud-Raught-2015

\mathbf{D}

Danielsen-Nielsen-2011 , Dart-Wells-2015 , de Hoog-Mann-2004 , Diner-Cristea-2015 , Dobbin-Giordano-2005 , Drissi-Boisvert-2015 , Dyer-Sobral-2010

 \mathbf{E}

Emanuele-Elledge-2011 , Emdal-Olsen-2015 , Ewing-Figeys-2007

 \mathbf{F}

Fenner-Prehn-2010 , Floyd-Pagliarini-2016 , Foerster-Ritter-2013 , Fogeron-Lange-2013 , Foster-Marshall-2013 , Freibaum-Taylor-2010

 \mathbf{G}

Gabriel-Baumgrass-2016, Galligan-Howley-2015, Gao-Reinberg-2012, Gautier-

 $\rm Hall-2009$, Giannone-Liu-2010 , Glatter-Gstaiger-2009 , Gloeckner-Ueffing-2007 , Goehler-Wanker-2004 , Golebiowski-Hay-2009 , Goudreault-Gingras-2009 , Grant-2010 , Greco-Cristea-2011 , Grossmann-Stelzl-2015 , Guarani-Harper-2014 , Gupta-Pelletier-2015

\mathbf{H}

Hanson-Clayton-2014 , Hauri-Gstaiger-2013 , Havrylov-Redowicz-2009 , Havugimana-Emili-2012 , Hayes-Urbé-2012 , Hegele-Stelzl-2012 A , Hegele-Stelzl-2012 B , Hein-Mann-2015 , Hill-Livingston-2014 , HUMANCYC , Humphries-Humphries-2009 , Hutchins-Peters-2010 , Huttlin-Gygi-2015

Ι

I2D-BIND-Fly2Human, I2D-BIND-Mouse2Human, I2D-BIND-Rat2Human, I2D-BIND-Worm2Human , I2D-BIND-Yeast2Human , I2D-BioGRID-Fly2Human , I2D-BioGRID-Mouse2Human, I2D-BioGRID-Rat2Human, I2D-BioGRID-Worm2Human, I2D-BioGRID-Yeast2Human, I2D-Chen-Pawson-2009-PiwiScreen-Mouse2Human, I2D-Formstecher-Daviet-2005-Embryo-Fly2Human, I2D-Giot-Rothbert-2003-Low-Fly2Human, I2D-INNATEDB-Mouse2Human, I2D-IntAct-Fly2Human, I2D-IntAct-Mouse2Human, I2D-IntAct-Rat2Human, I2D-IntAct-Worm2Human, I2D-IntAct-Yeast2Human, I2D-Krogan-Greenblatt-2006-Core-Yeast2Human, I2D-Krogan-Greenblatt-2006-NonCore-Yeast2Human, I2D-Li-Vidal-2004-CORE-1-Worm2Human, I2D-Li-Vidal-2004-non-core-Worm2Human, I2D-Manual-Mouse2Human , I2D-Manual-Rat2Human , I2D-MGI-Mouse2Human . I2D-MINT-Fly2Human, I2D-MINT-Mouse2Human, I2D-MINT-Rat2Human, I2D-MINT-Worm2Human, I2D-MINT-Yeast2Human, I2D-Ptacek-Snyder-2005-Yeast2Human, I2D-Tarassov-PCA-Yeast2Human, I2D-Tewari-Vidal-2004-TGFb-Worm2Human, I2D-vonMering-Bork-2002-High-Yeast2Human, I2D-vonMering-Bork-2002-Low-Yeast2Human, I2D-vonMering-Bork-2002-Medium-Yeast2Human, I2D-Wang-Orkin-2006-EScmplx-Mouse2Human, I2D-Wang-Orkin-2006-EScmplxlow-Mouse2Human, I2D-Yu-Vidal-2008-GoldStd-Yeast2Human, IMID, Ingham-Pawson-2005, Innocenti-Brown-2011, INTERPRO, IREF-BIND, IREF-BIOGRID, IREF-DIP, IREF-HPRD, IREF-INTACT, IREF-MATRIXDB, IREF-MPPI, IREF-PUBMED, IREF-SMALL-SCALE-STUDIES, IREF-SMALL-SCALE-STUDIES

J

Jeronimo-Coulombe-2007 , Jin-Pawson-2004 , Johnson-Kerner-Wichterle-2015 , Johnson-Shoemaker-2003 , Jones-MacBeath-2006 , Joshi-Cristea-2013 , Jäger-Krogan-2011

\mathbf{K}

Kahle-Zoghbi-2011, Kaltenbach-Hughes-2007, Katsogiannou-Rocchi-2014, Kim-Gygi-2011, Kim-Major-2015, Kneissl-Grummt-2003, Koch-Hermeking-2007, Kotlyar-Jurisica-2015, Kristensen-Foster-2012, Kärblane-Sarmiento-2015, Kırlı-

Görlich-2015

\mathbf{L}

 $Lambert-Gingras-2015\ ,\ Lamoliatte-Thibault-2014\ ,\ Lau-Ronai-2012\ ,\ Lee-Songyang-2011\ ,\ Lehner-Sanderson-2004\ A\ ,\ Lehner-Sanderson-2004\ B\ ,\ Leng-Wang-2014\ ,\ Leung-Jones-2014\ ,\ Li-Chen-2015\ ,\ Li-Dorf-2011\ A\ ,\ Li-Dorf-2011\ B\ ,\ Li-Dorf-2014\ ,\ Li-Haura-2013\ ,\ Lim-Zoghbi-2006\ ,\ Lin-Smith-2010\ ,\ Lipp-Guthrie-2015\ ,\ Liu-Wang-2012\ ,\ Llères-Lamond-2010\ ,\ Loch-Strickler-2012\ ,\ Low-Heck-2014\ ,\ Lu-Zhang-2013\ ,\ Luo-Elledge-2009$

\mathbf{M}

Mak-Moffat-2010 , Mallon-McKay-2013 , Malovannaya-Qin-2010 , Markson-Sanderson-2009 , Maréchal-Zou-2014 , Matsumoto-Nakayama-2005 , McCracken-Blencowe-2005 , McFarland-Nussbaum-2008 , Meek-Piwnica-Worms-2004 , Milev-Mouland-2012 , Miyamoto-Sato-Yanagawa-2010 , Murakawa-Landthaler-2015

\mathbf{N}

Nakayama-Ohara-2002 , Nakayasu-Adkins-2013 , Napolitano-Meroni-2011 , Narayan-Bennett-2012 , Nathan-Goldberg-2013 , NCI_NATURE , Neganova-Lako-2011 , Newman-Keating-2003 , Nicholson-Hupp-2014 , Noble-Diehl-2008

O

Oliviero-Cagney-2015 , Olma-Pintard-2009 , Oláh-Ovádi-2011 , Oshikawa-Nakayama-2012 , Ouyang-Gill-2009

\mathbf{P}

Panigrahi-Pati-2012 , Papp-Lamia-2015 , Perez-Hernandez-Yáñez-Mó-2013 , Perou-Botstein-1999 , Perou-Botstein-2000 , Persaud-Rotin-2009 , Petschnigg-Stagljar-2014 , PFAM , Phillips-Corn-2013 , Pichlmair-Superti-Furga-2011 , Pichlmair-Superti-Furga-2012 , Pilot-Storck-Goillot-2010 , Povlsen-Choudhary-2012

${f R}$

Ramachandran-LaBaer-2004 , Raman-Harper-2015 , Ramaswamy-Golub-2001 , Ravasi-Hayashizaki-2010 , REACTOME , Reinke-Keating-2013 , Reyniers-Taymans-2014 , Richter-Chrzanowska-Lightowlers-2010 , Rieger-Chu-2004 , Rolland-Vidal-2014 , Rosenwald-Staudt-2001 , Roth-Zlotnik-2006 , Roux-Burke-2012 , Rowbotham-Mermoud-2011 , Roy-Pardo-2014 , Roy-Parent-2013 , Rual-Vidal-2005 A , Rual-Vidal-2005 B

Ç

Sang-Jackson-2011 , Sato-Conaway-2004 , Schadt-Shoemaker-2004 , Scholz-Taylor-2016 , Singh-Moore-2012 , Smirnov-Cheung-2009 , So-Colwill-2015 , Soler-López-Aloy-2011 , Sowa-Harper-2009 , Stehling-Lill-2012 , Stehling-Lill-2013 , Stelzl-Wanker-2005 , Stes-Gevaert-2014 , Stuart-Kim-2003 , Suter-Wanker-2013

\mathbf{T}

\mathbf{T}

Taipale-Lindquist-2012 , Taipale-Lindquist-2014 , Takahashi-Conaway-2011 , Tarallo-Weisz-2011 , Tatham-Hay-2011 , Teixeira-Gomes-2010 , Thalappilly-Dusetti-2008 , Thompson-Luchansky-2014 , Tong-Moran-2014 , Toyoshima-Grandori-2012 , Tsai-Cristea-2012

\mathbf{U}

Udeshi-Carr-2012

\mathbf{V}

van Wijk-Timmers-2009 , Vandamme-Angrand-2011 , Varjosalo-Gstaiger-2013 , Varjosalo-Superti-Furga-2013 , Venkatesan-Vidal-2009 , Vermeulen-Mann-2010 , Vinayagam-Wanker-2011 , Virok-Fülöp-2011 , Vizeacoumar-Moffat-2013

\mathbf{W}

Wagner-Choudhary-2011 , Wallach-Kramer-2013 , Wan-Emili-2015 , Wang-Balch-2006 , Wang-Cheung-2015 , Wang-He-2008 , Wang-Maris-2006 , Wang-Xu-2015 , Wang-Yang-2011 , Weimann-Stelzl-2013 A , Weimann-Stelzl-2013 B , Weimann-Meister-2009 , Wen-Wu-2014 , Whisenant-Salomon-2015 , Wilker-Yaffe-2007 , Willingham-Muchowski-2003 , Witt-Labeit-2008 , Wong-O'Bryan-2012 , Woods-Monteiro-2012 , Woodsmith-Sanderson-2012 , Wu-Garvey-2007 , Wu-Li-2007 , Wu-Ma-2012 , Wu-Stein-2010 , Wu-Stein-2010

\mathbf{X}

Xiao-Lefkowitz-2007, Xie-Cong-2013, Xie-Green-2012, Xu-Ye-2012

Y

Yang-Chen-2010, Yatim-Benkirane-2012, Yu-Chow-2013, Yu-Vidal-2011

${f Z}$

Zanon-Pichler-2013 , Zhang-Shang-2006 , Zhang-Zou-2011 , Zhao-Krug-2005 , Zhao-Yang-2011 , Zhou-Conrads-2004 , Zhou-Hanemann-2016

Genes

Gene	Description	Rank
LTA4H	leukotriene A4 hydrolase [Source:HGNC Symbol;Acc:HGNC:6710]	N/A
LMCD1	LIM and cysteine rich domains 1 [Source:HGNC Symbol;Acc:HGNC:6633]	N/A
MTUS1	microtubule associated tumor suppressor 1 [Source:HGNC Symbol;Acc: HGNC:29789]	N/A
HTR2B	5-hydroxytryptamine receptor 2B [Source:HGNC Symbol;Acc:HGNC: 5294]	N/A
FXR1	FMR1 autosomal homolog 1 [Source:HGNC Symbol;Acc:HGNC:4023]	N/A
PRAME	preferentially expressed antigen in melanoma [Source:HGNC Symbol;Acc: HGNC:9336]	N/A
ID2	inhibitor of DNA binding 2, HLH protein [Source:HGNC Symbol;Acc: HGNC:5361]	N/A
CDH1	cadherin 1 [Source:HGNC Symbol;Acc:HGNC:1748]	N/A
EIF1B	eukaryotic translation initiation factor 1B [Source:HGNC Symbol;Acc: HGNC:30792]	N/A
ECM1	extracellular matrix protein 1 [Source:HGNC Symbol;Acc:HGNC:3153]	N/A
RAB31	RAB31, member RAS oncogene family [Source:HGNC Symbol;Acc: HGNC:9771]	N/A
ROBO1	roundabout guidance receptor 1 [Source:HGNC Symbol;Acc:HGNC:10249]	N/A
SATB1	SATB homeobox 1 [Source:HGNC Symbol;Acc:HGNC:10541]	N/A
SLK	STE20 like kinase [Source:HGNC Symbol;Acc:HGNC:11088]	1
MIB1	mindbomb E3 ubiquitin protein ligase 1 [Source:HGNC Symbol;Acc: HGNC:21086]	2
USP8	ubiquitin specific peptidase 8 [Source:HGNC Symbol;Acc:HGNC:12631]	3
DAB2	DAB2, clathrin adaptor protein [Source:HGNC Symbol;Acc:HGNC:2662]	4
MFGE8	milk fat globule-EGF factor 8 protein [Source:HGNC Symbol;Acc:HGNC: 7036]	5
TMEFF1	transmembrane protein with EGF like and two follistatin like domains 1 [Source:HGNC Symbol;Acc:HGNC:11866]	6
GOLGA4	golgin A4 [Source:HGNC Symbol;Acc:HGNC:4427]	7
MEST	mesoderm specific transcript [Source:HGNC Symbol;Acc:HGNC:7028]	8
MOB3A	MOB kinase activator 3A [Source:HGNC Symbol;Acc:HGNC:29802]	9
SHB	SH2 domain containing adaptor protein B [Source:HGNC Symbol;Acc: HGNC:10838]	10

Gene	Description	Rank
MTHFD2	methylenetetrahydrofolate dehydrogenase (NADP+ dependent) 2, methenyltetrahydrofolate cyclohydrolase [Source:HGNC Symbol;Acc: HGNC:7434]	11
THBS1	thrombospondin 1 [Source:HGNC Symbol;Acc:HGNC:11785]	12
LHFPL2	lipoma HMGIC fusion partner-like 2 [Source:HGNC Symbol;Acc:HGNC: 6588]	13
DYNC1LI2	dynein cytoplasmic 1 light intermediate chain 2 [Source:HGNC Symbol; Acc:HGNC:2966]	14
FOXG1	forkhead box G1 [Source:HGNC Symbol;Acc:HGNC:3811]	15
VDR	vitamin D (1,25- dihydroxyvitamin D3) receptor [Source:HGNC Symbol; Acc:HGNC:12679]	16
STK26	serine/threonine protein kinase 26 [Source:HGNC Symbol;Acc:HGNC: 18174]	17
DTX4	deltex 4, E3 ubiquitin ligase [Source:HGNC Symbol;Acc:HGNC:29151]	18
TLL2	tolloid like 2 [Source:HGNC Symbol;Acc:HGNC:11844]	19
DSP	desmoplakin [Source:HGNC Symbol;Acc:HGNC:3052]	20

Networks

Co-expression	100. $00%$
Chen-Brown-2002	34.84%
Gene expression patterns in human liver cancers. Chen et al (2002). Mol Biol Cell	
Co-expression with 282,241 interactions from supplementary material	
Wang-Cheung-2015	27.17%
Genetic variation in insulin-induced kinase signaling. Wang et al (2015). Mol Syst Biol	
Co-expression with 411,047 interactions from GEO	
Burington-Shaughnessy-2008	19.69%
Tumor cell gene expression changes following short-term in vivo exposure to single agent chemotherapeutics are related to survival in multiple myeloma. Burington et al (2008) . Clin Cancer Res	
Co-expression with 290,538 interactions from GEO	
Rieger-Chu-2004	9.90%
Toxicity from radiation therapy associated with abnormal transcriptional responses to DNA damage. Rieger et al (2004). $Proc$ $Nath\ Acad\ Sci\ U\ S\ A$	
Co-expression with 259,974 interactions from GEO	
Dobbin-Giordano-2005	8.23%
Interlaboratory comparability study of cancer gene expression analysis using oligonucleotide microarrays. Dobbin et al (2005) . Clin Cancer Res	
Co-expression with 444,931 interactions from GEO	
Wu-Garvey-2007	
The effect of insulin on expression of genes and biochemical pathways in human skeletal muscle. Wu et al (2007). Endocrine	
Co-expression with 267,109 interactions from GEO	