

While parsing you can click empty space and and drag to zoom
Could not parse GO ID from line: 'Term'. Line will be skipped.

interpretation of the coordinate axes?



Hide/show dispensable GO terms

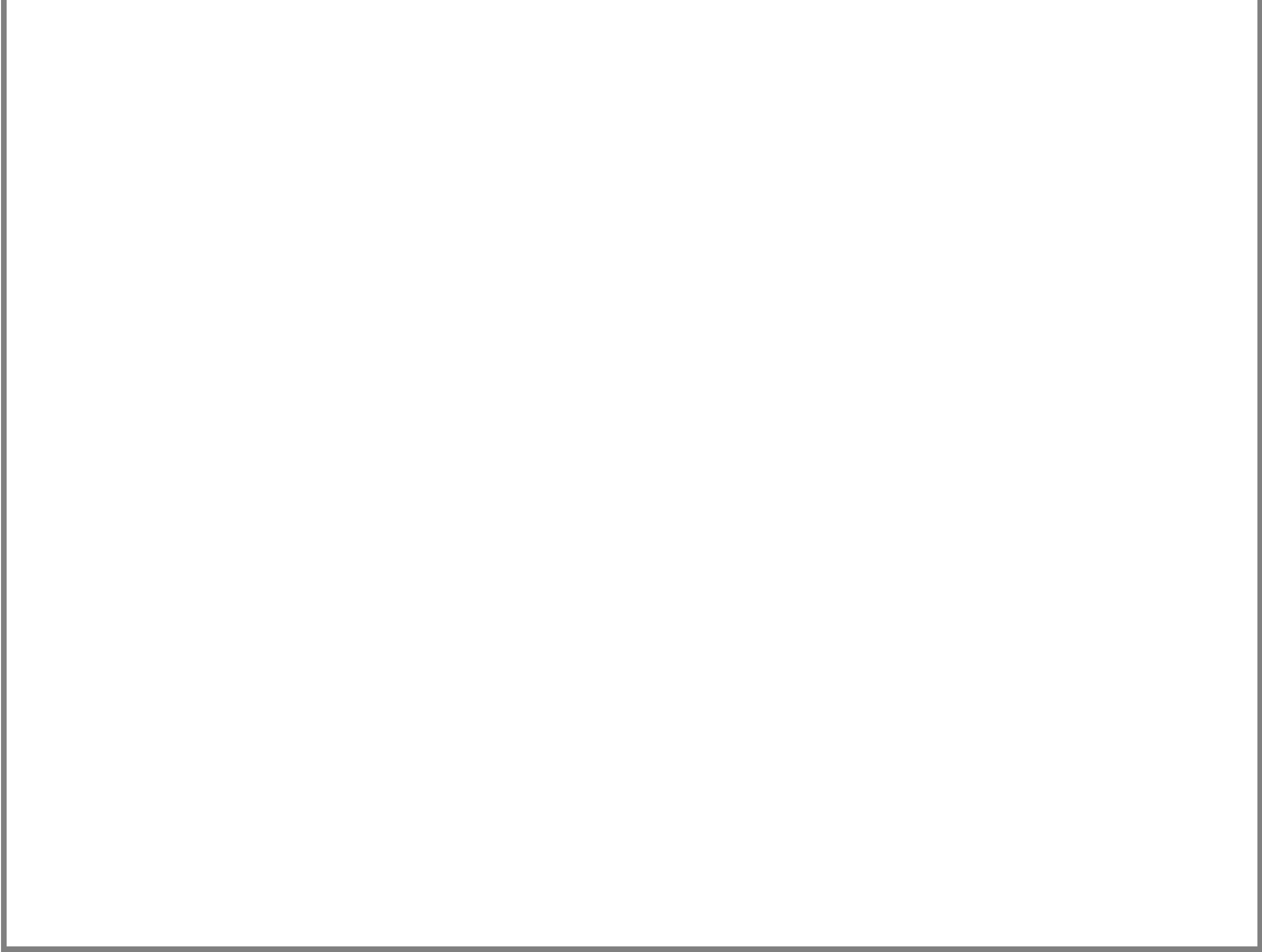


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term ID	description	frequency	pin?	uniqueness	dispensability
GO:0001649	osteoblast differentiation	0.013 %		0.82	0.00
GO:0033280	response to vitamin D	0.001 %		0.79	0.00
GO:0051028	mRNA transport	0.053 %		0.75	0.00
GO:0030490	maturation of SSU-rRNA	0.001 %		0.70	0.01
GO:0042359	vitamin D metabolic process	0.001 %		0.69	0.06
GO:0042369	<i>vitamin D catabolic process</i>	0.000 %		0.69	0.86
GO:0055114	oxidation-reduction process	15.044 %		0.83	0.08
GO:0010467	gene expression	16.702 %		0.77	0.11
GO:0006810	transport	17.382 %		0.79	0.31
GO:0035754	B cell chemotaxis	0.000 %		0.68	0.34
GO:0008380	RNA splicing	0.097 %		0.66	0.38
GO:0006397	<i>mRNA processing</i>	0.373 %		0.65	0.56
GO:1900182	positive regulation of protein localization to nucleus	0.005 %		0.61	0.44
GO:0006956	<i>complement activation</i>	0.069 %		0.54	0.66
GO:0045727	<i>positive regulation of translation</i>	0.047 %		0.60	0.62



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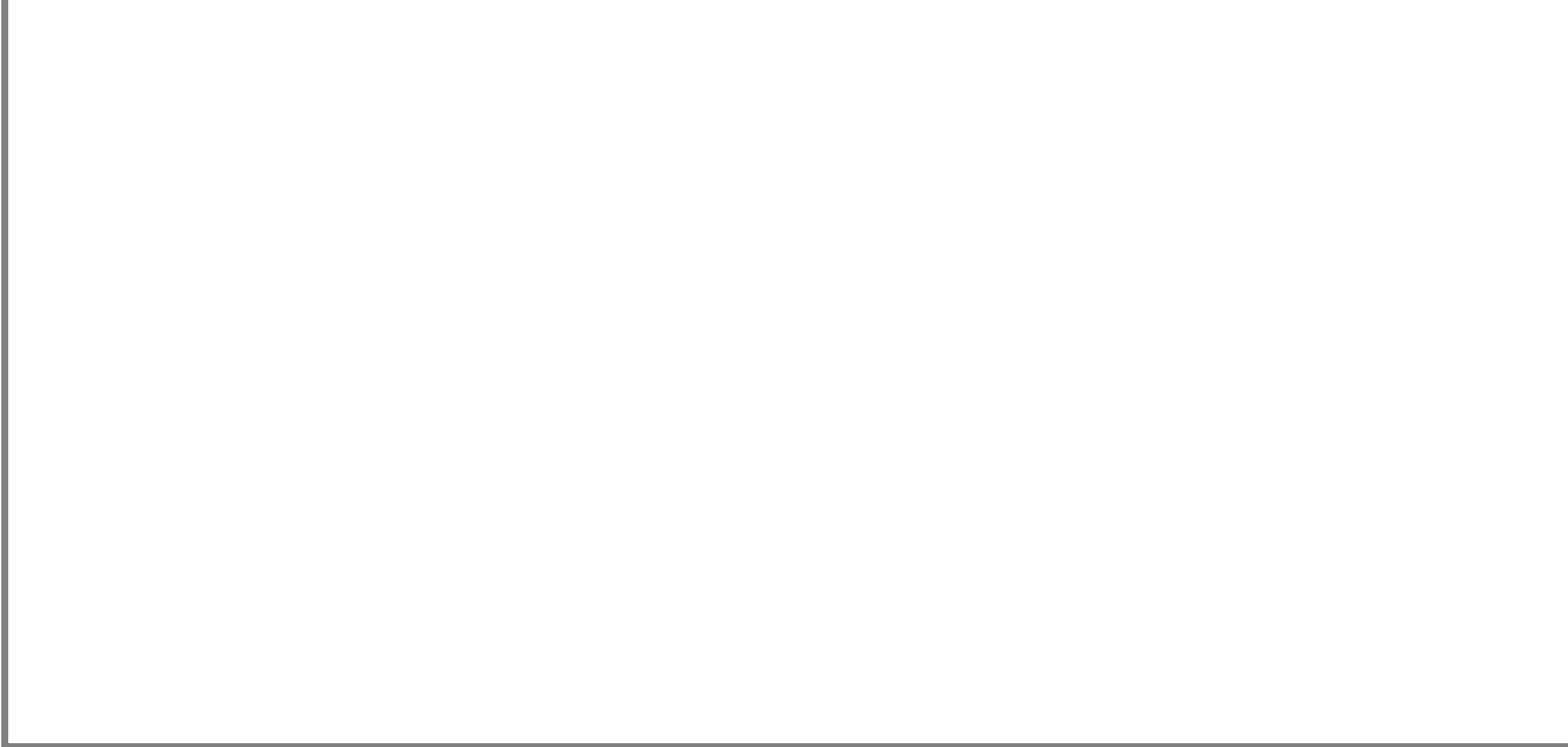
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term ID	description	frequency	pin?	uniqueness	dispensability
GO:0016235	aggresome	0.003 %		0.51	0.00
GO:0070062	extracellular exosome	0.300 %		0.43	0.00
GO:0005886	plasma membrane	13.934 %		0.58	0.03
GO:0048471	perinuclear region of cytoplasm	0.050 %		0.51	0.05
GO:0005654	nucleoplasm	0.221 %		0.38	0.37

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term ID	description	frequency	pin?	uniqueness	dispensability
GO:0003939	L-iditol 2-dehydrogenase activity	0.003 %		0.70	0.00
GO:0042803	protein homodimerization activity	0.122 %		0.73	0.00
GO:0016491	oxidoreductase activity	14.657 %		0.79	0.02
GO:0008144	drug binding	0.260 %		0.73	0.04
GO:0044822	poly(A) RNA binding	0.135 %		0.72	0.04
GO:0005506	iron ion binding	2.467 %		0.70	0.05
GO:0008403	25-hydroxycholecalciferol-24-hydroxylase activity	0.000 %		0.54	0.13
 GO:0030342	1-alpha,25-dihydroxyvitamin D3 24-hydroxylase activity	0.000 %		0.54	0.95
GO:0005525	GTP binding	1.692 %		0.69	0.19

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Frequent keywords within your set of GO terms:

localisation anabolic modifications single-organism calciol d2 d3 fluid-filled triacylglycerols subcellular macromolecular functioning transform
compound c-24 enclosed inclusion membrane-enclosed living plastids insoluble ch-oh individual weight repetition organic biopolymer
localization solvent non-encoded sparingly 25-hydroxycholecalciferol guided substance aggregated vacuoles dimerization calciferol ribosomes
self-propelled multiple location fat-soluble vitamins encapsulating steroids 24-hydroxylase low protoplast 25-hydroxyvitamin
localizer protein-coding delta-5 alcohols substances ergocalciferol solvents subcomponent immunogenic macromolecule
outermost organelle cholecalciferol housed ortho)phosphate bounded cellular macromolecules cyclopentanoperhydrophenanthrene oligophosphate

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