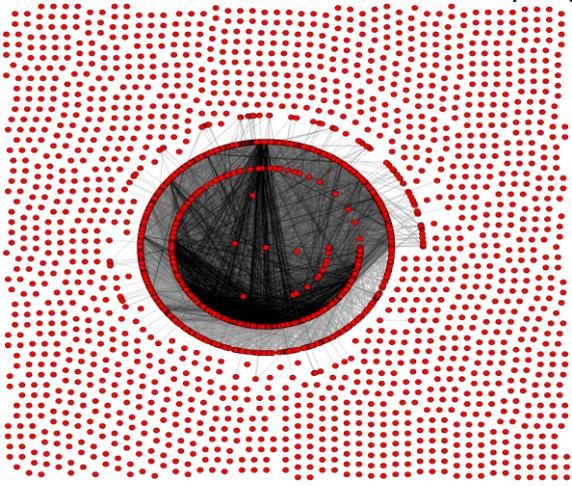
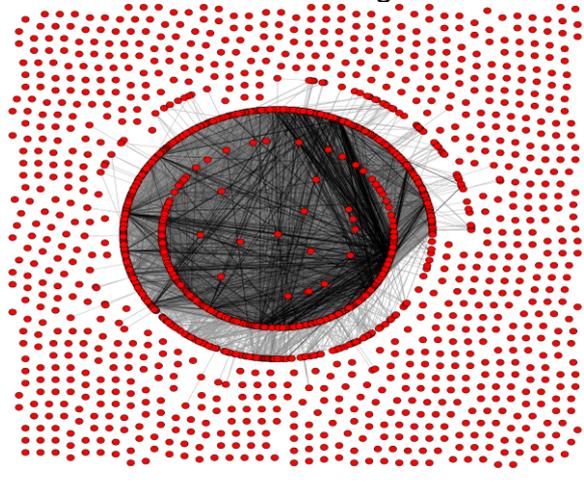


Contact networks for GO-slim terms

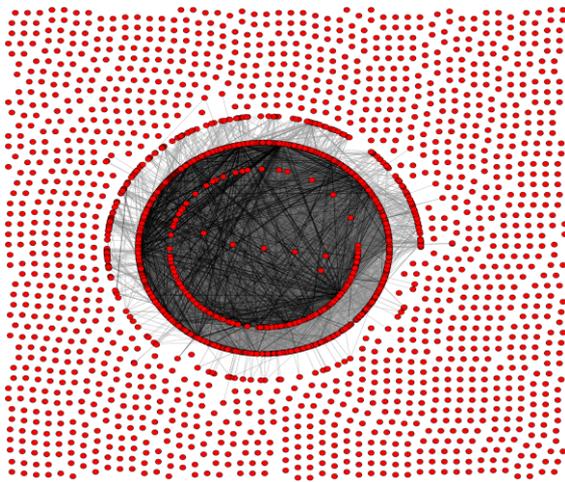
The figures below show the contact networks (frequency > 5) for each of the Go-slim terms. The number of links per gene is shown below each figure.



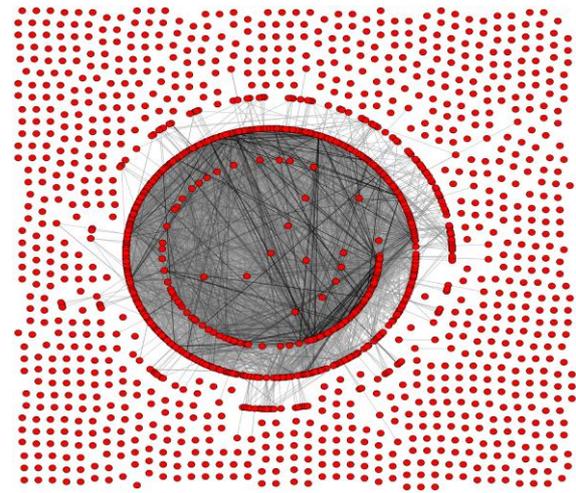
cytoplasm 49763 / 3401



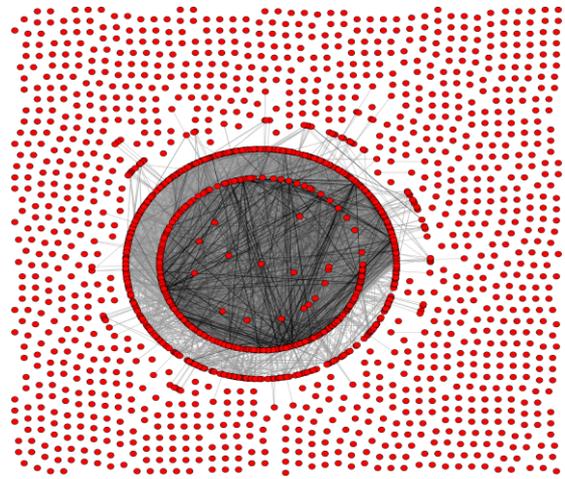
nucleus 17566 / 1852



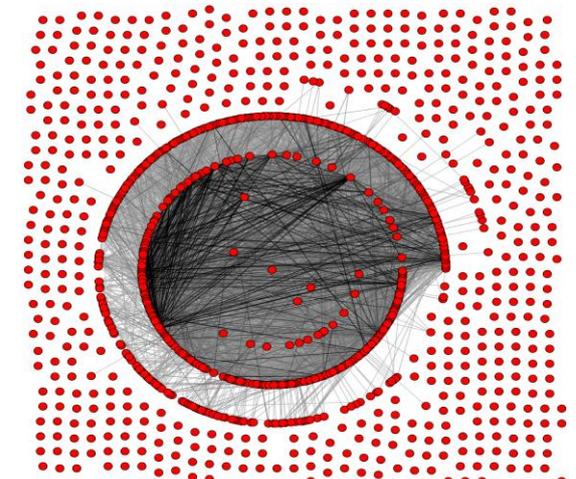
molecular function 22940 / 2734



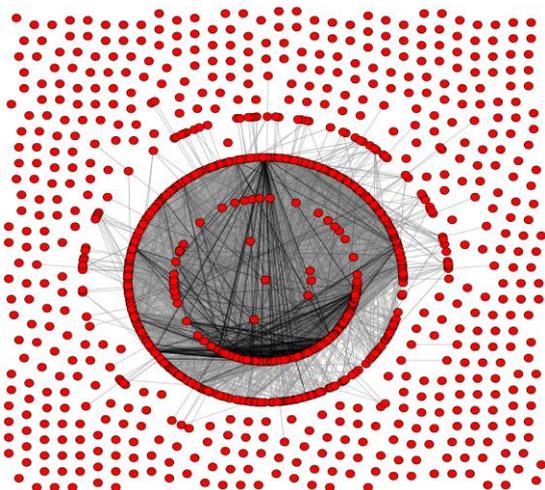
cellular component 5091 / 1548



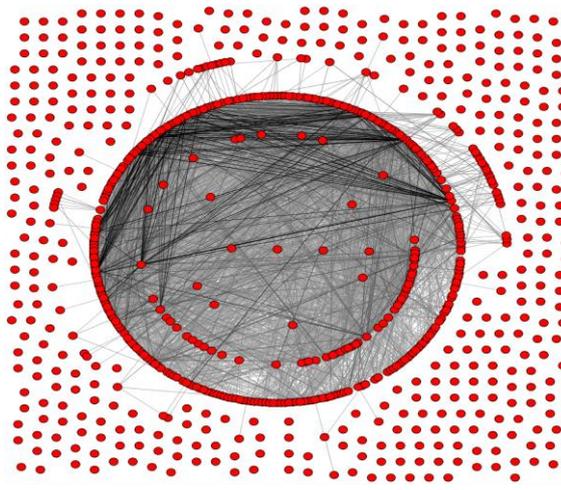
biological process 9233 / 1992



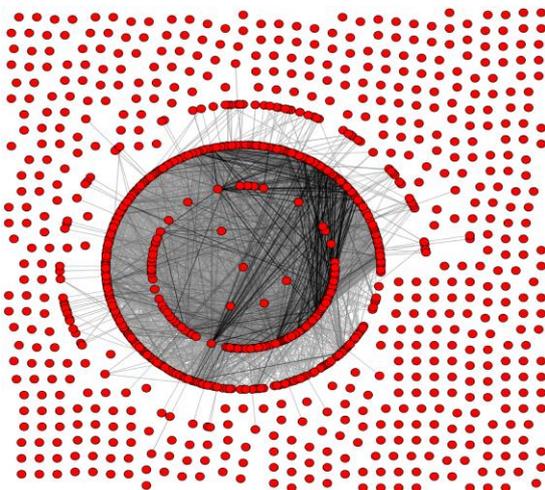
membrane 6618 / 1114



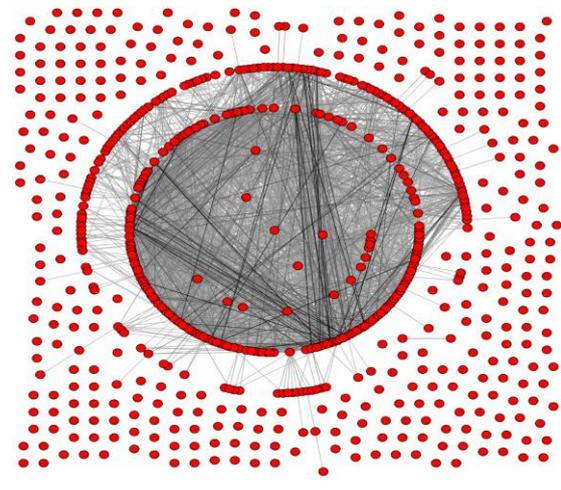
RNA metabolic process 5676 / 1063



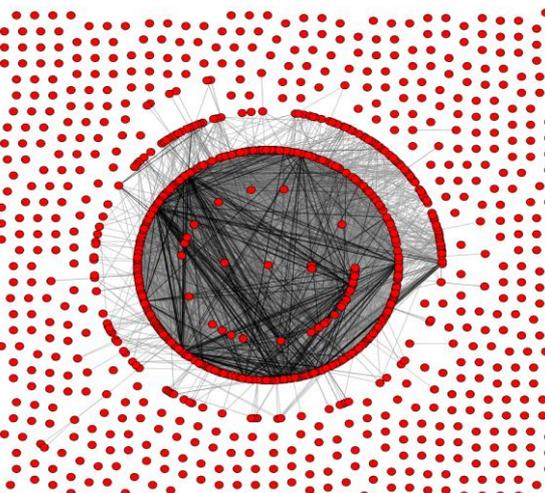
hydrolase activity 3688 / 779



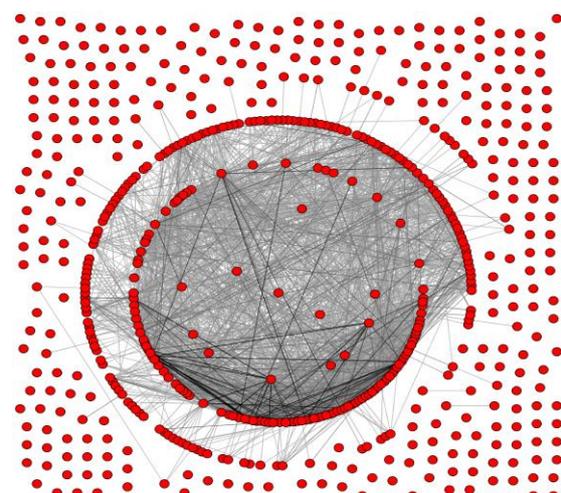
mitochondrion 4476 / 1039



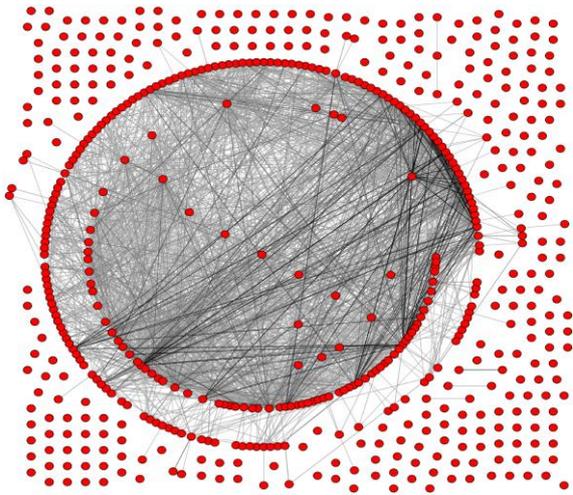
transferase activity 2232 / 662



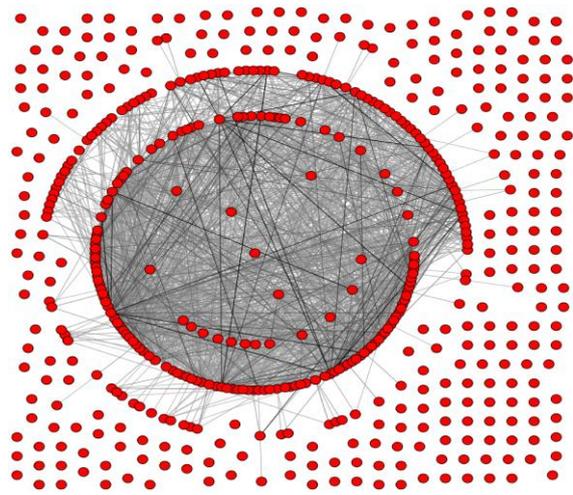
transport 5099 / 1010



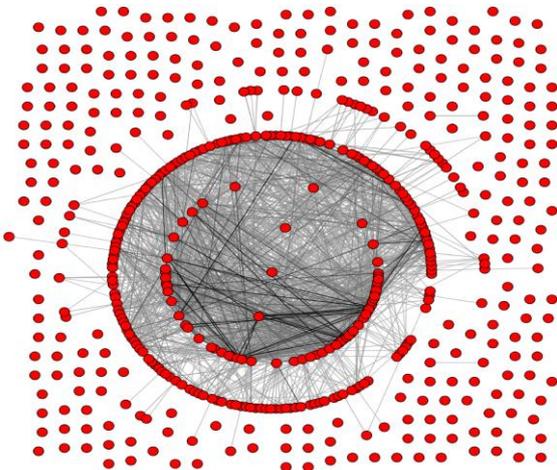
transcription 2047 / 596



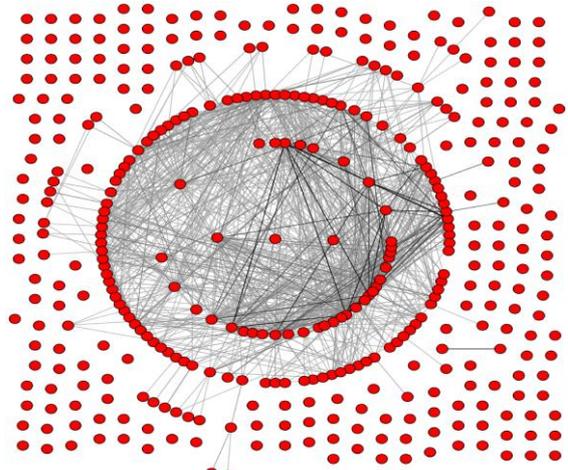
response to stress 1785 / 561



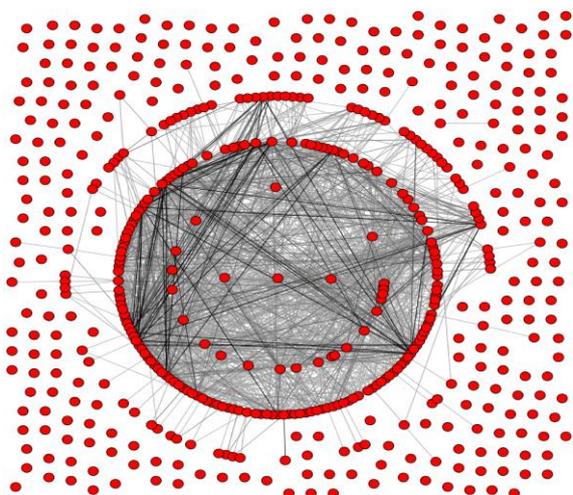
cell cycle 1870 / 521



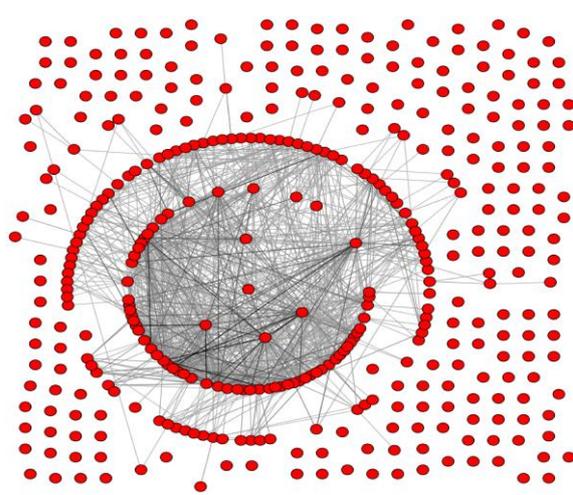
protein modification process 1561 / 537



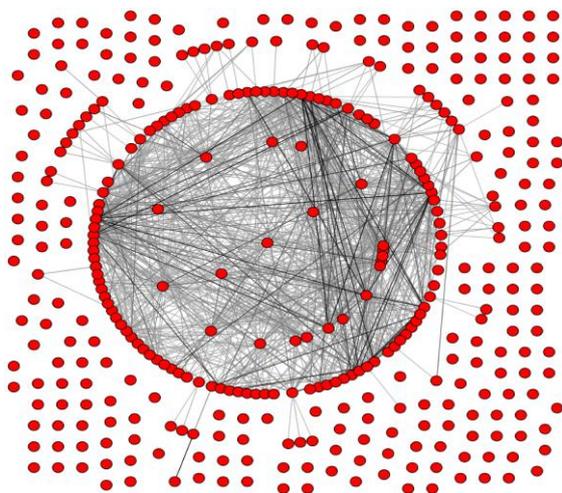
DNA metabolic process 839 / 388



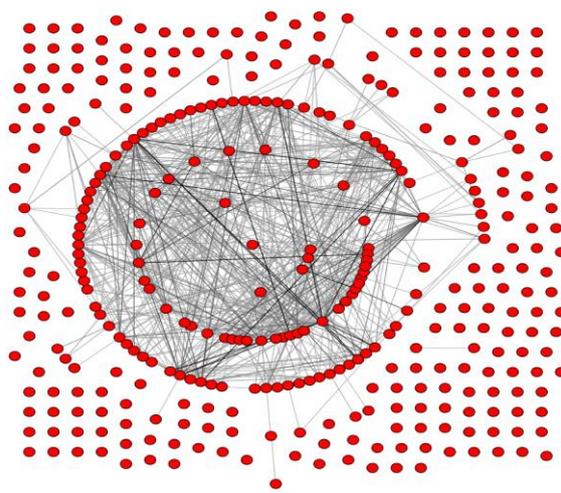
protein binding 1418 / 527



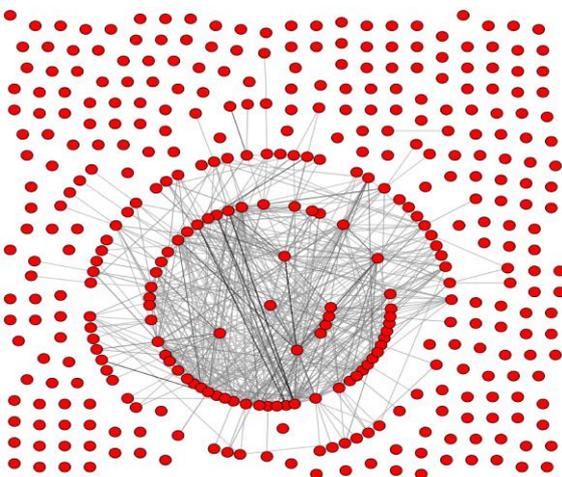
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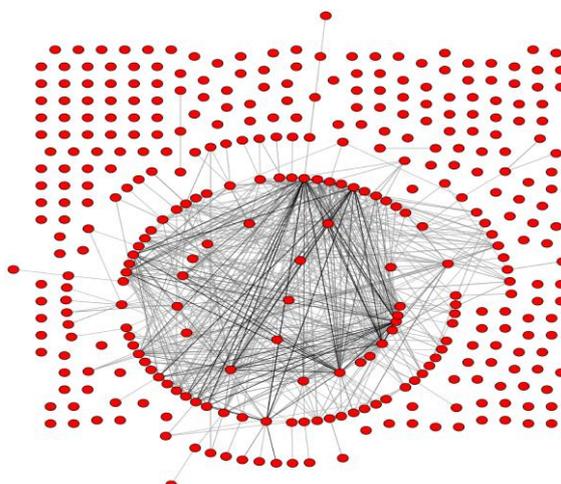
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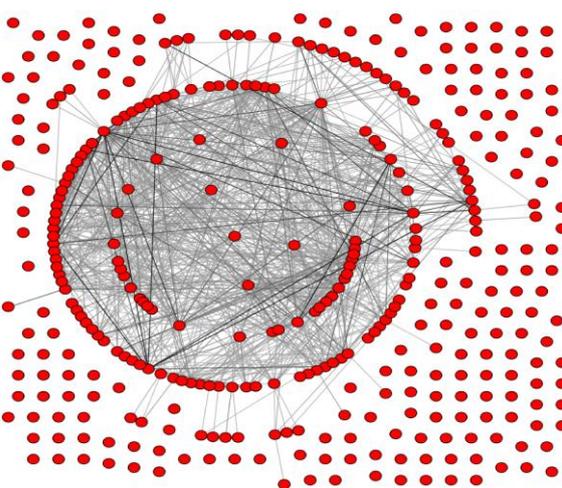
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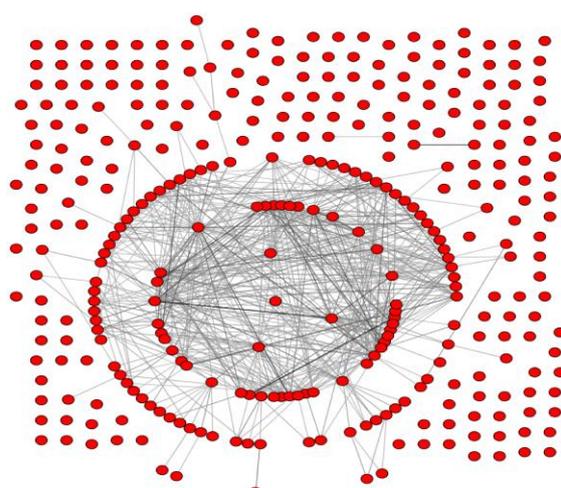
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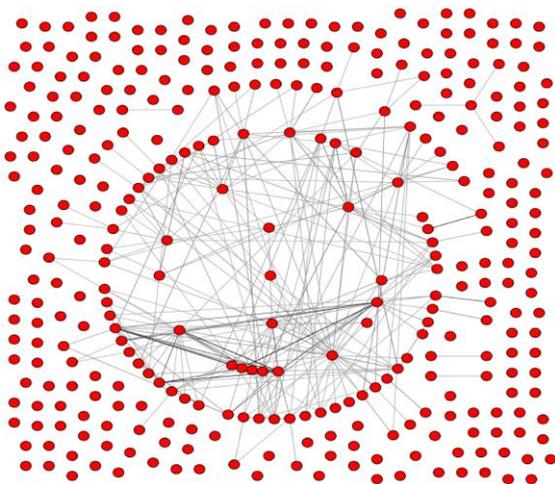
vesicle-mediated transport 511 / 347



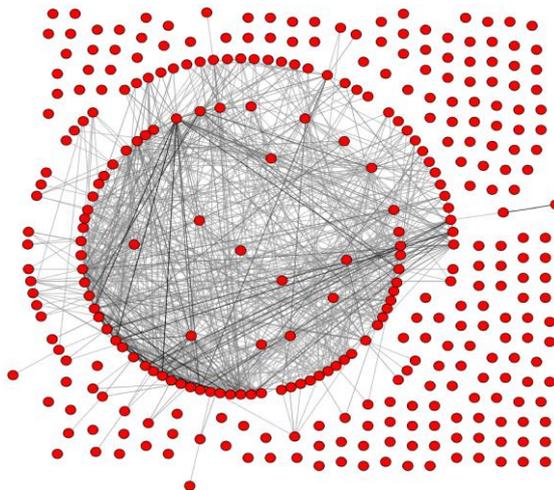
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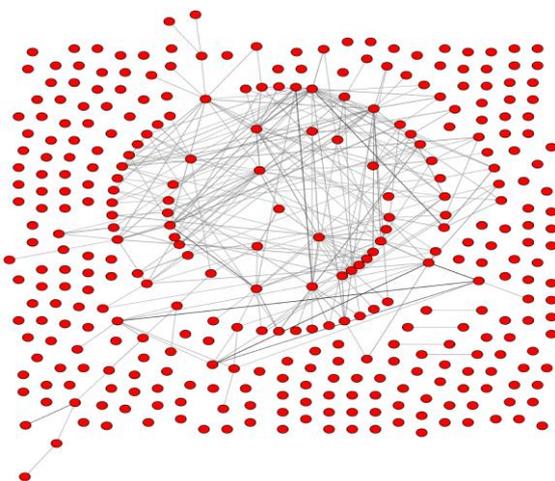
DNA binding 510 / 334



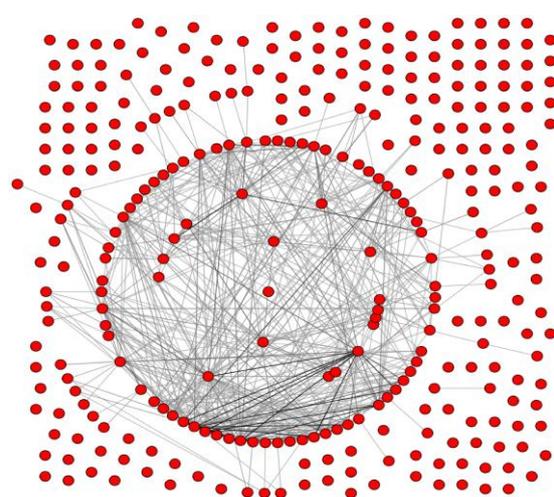
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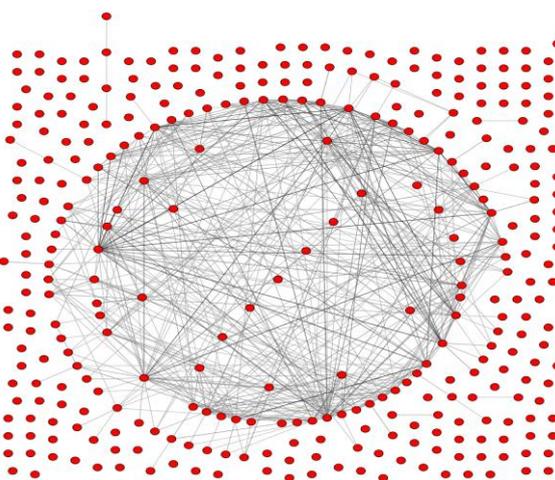
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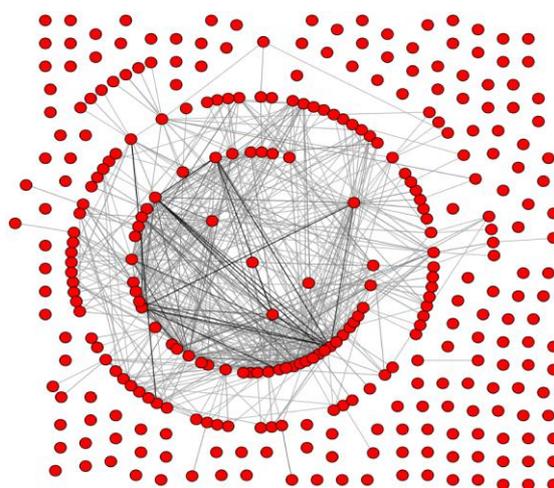
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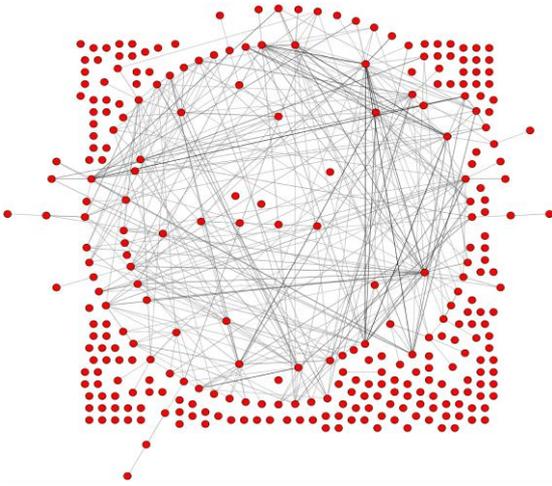
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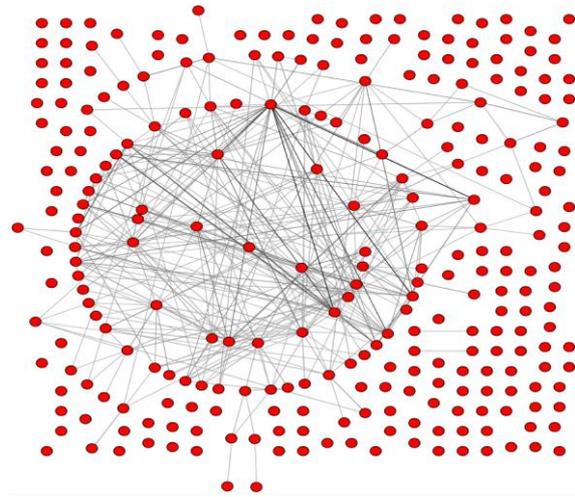
ribosome biogenesis 471 / 330



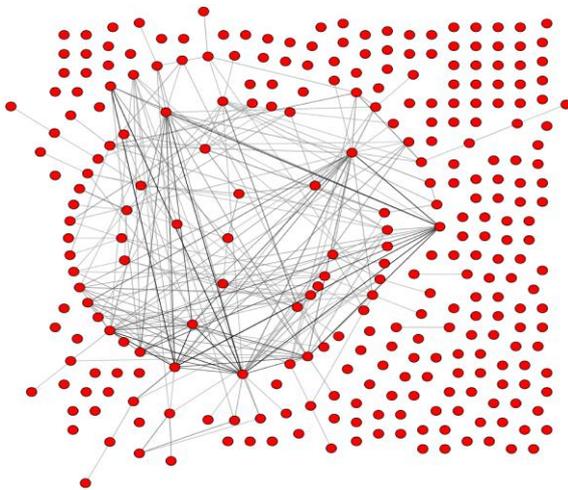
transcription regulator activity 472 / 312



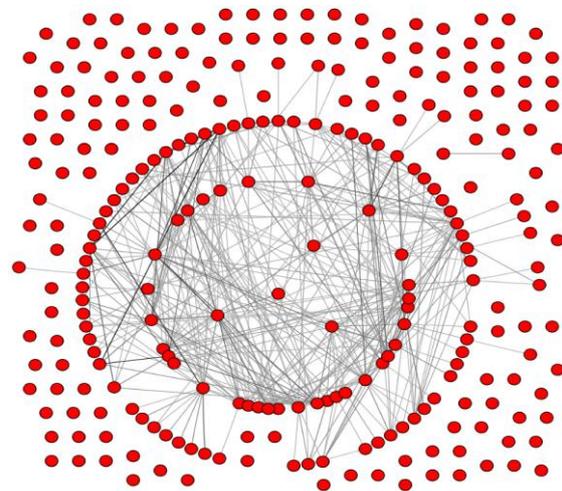
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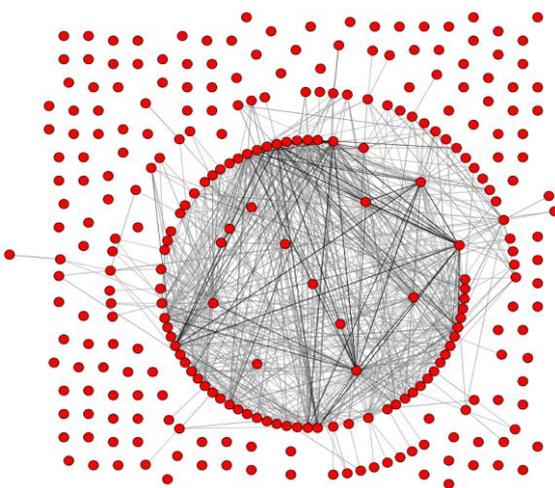
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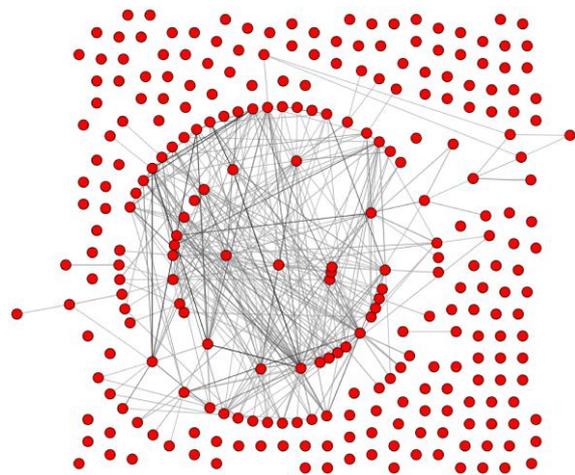
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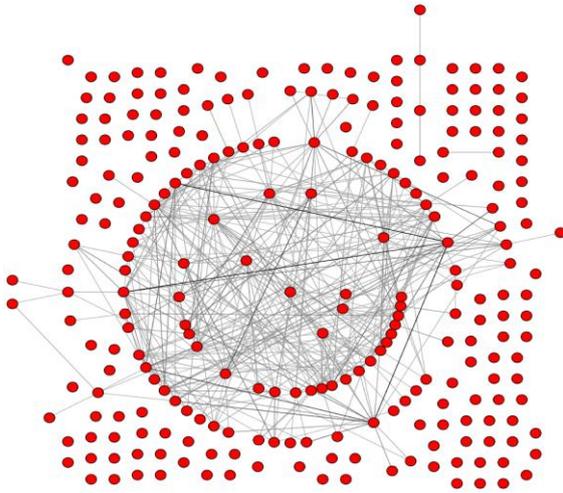
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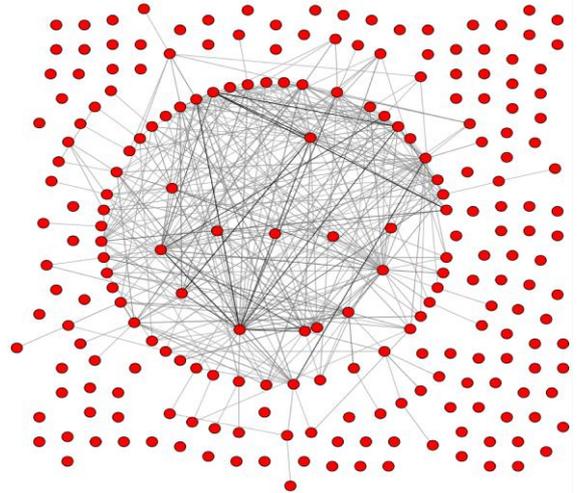
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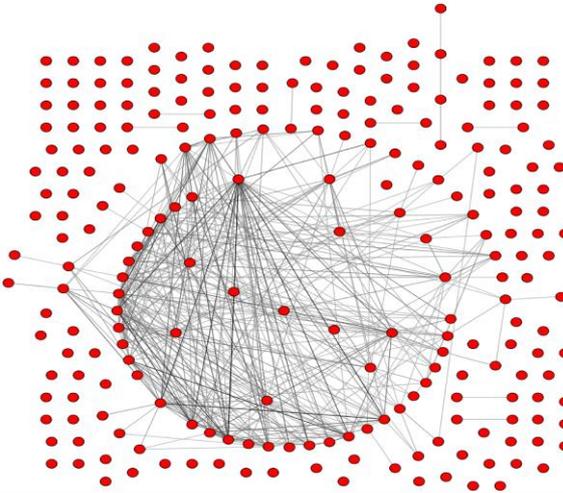
oxidoreductase activity 303 / 265



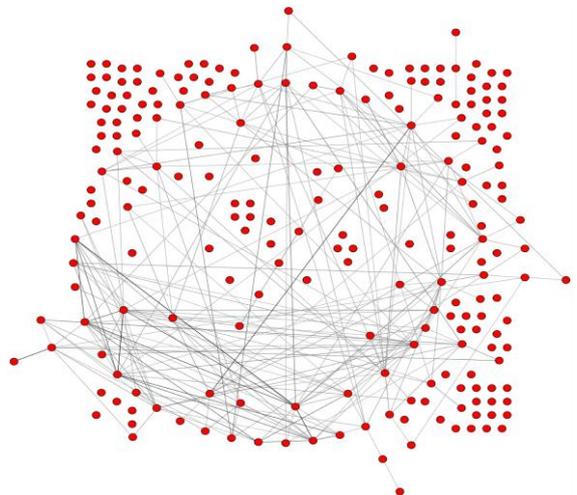
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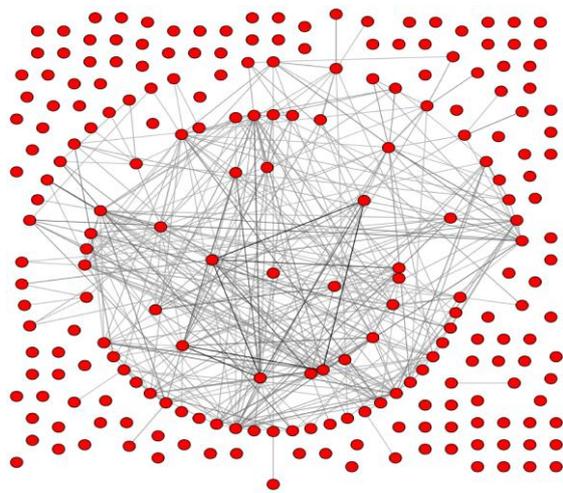
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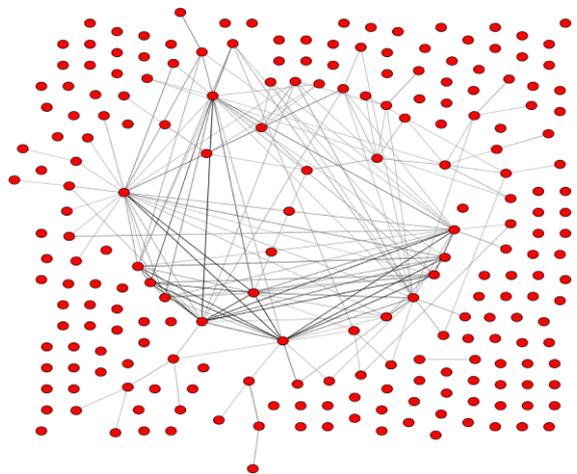
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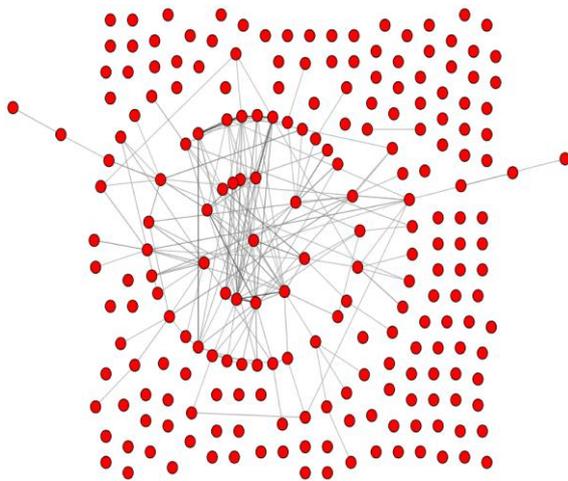
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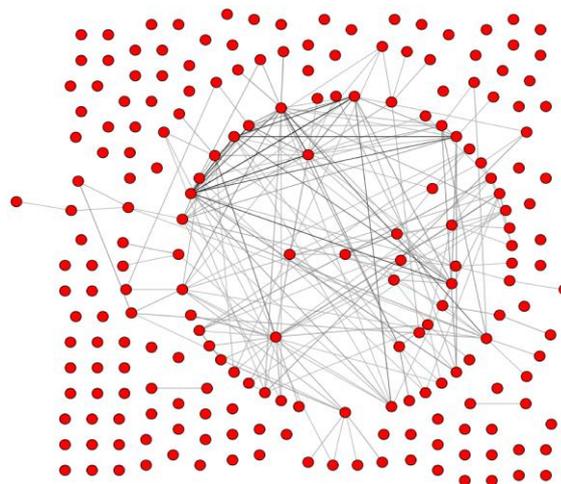
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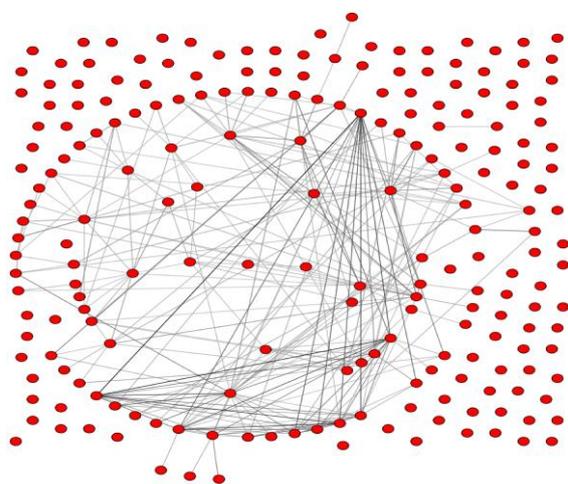
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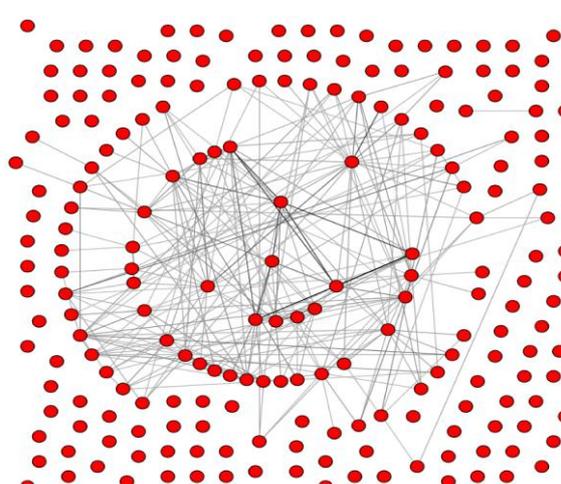
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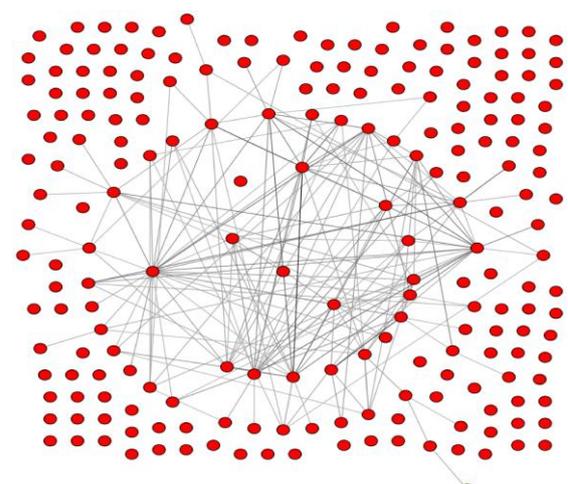
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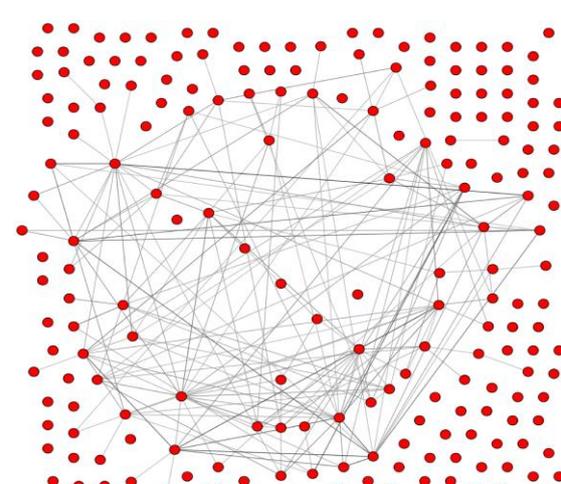
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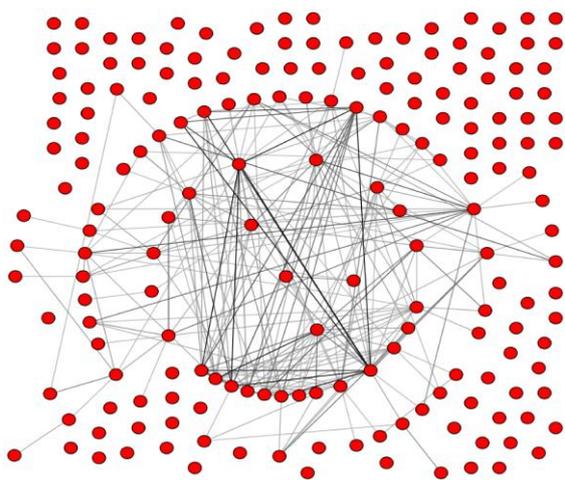
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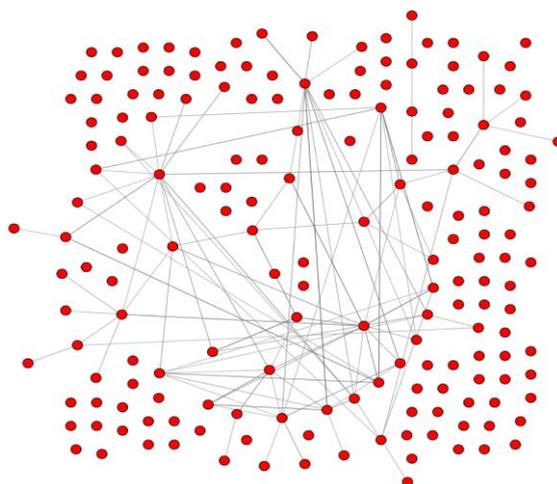
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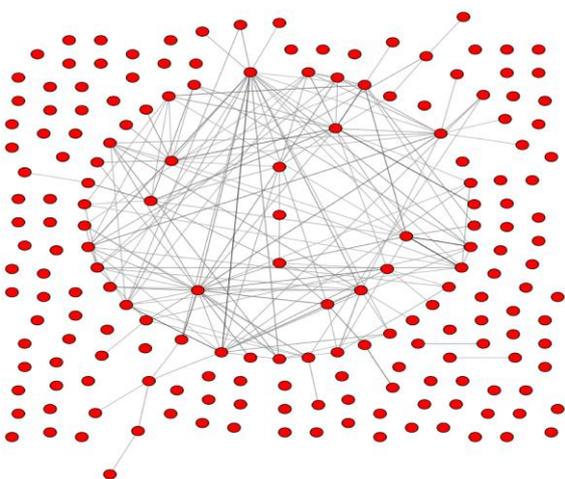
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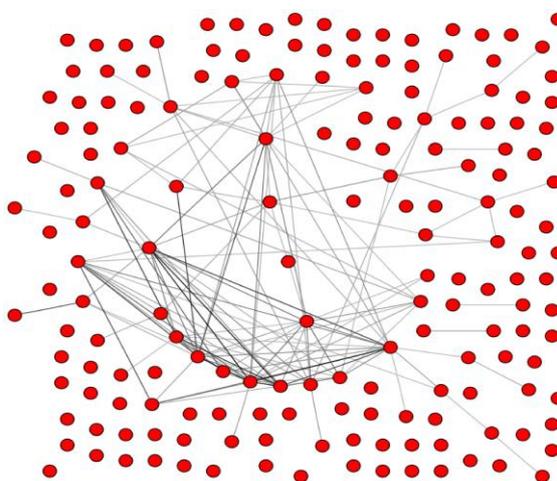
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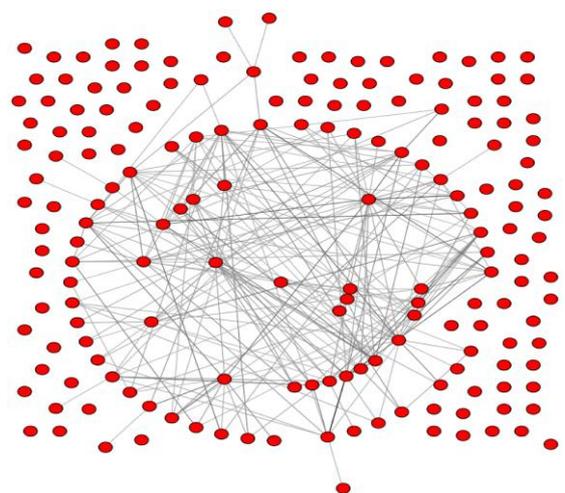
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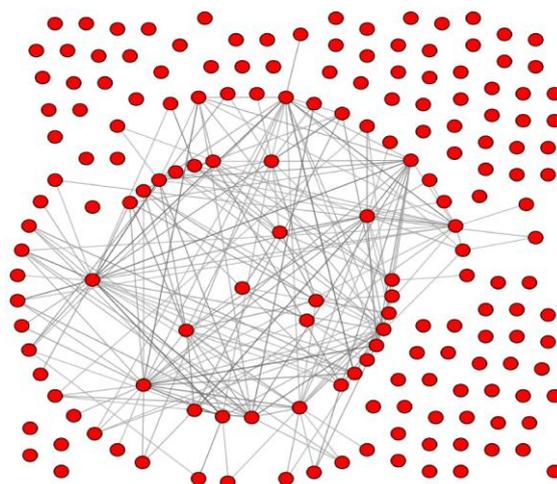
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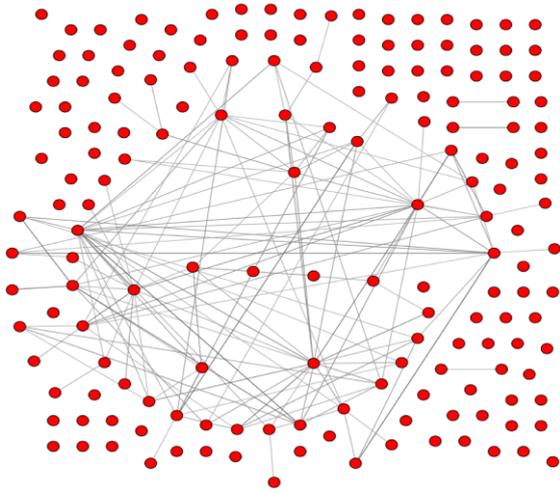
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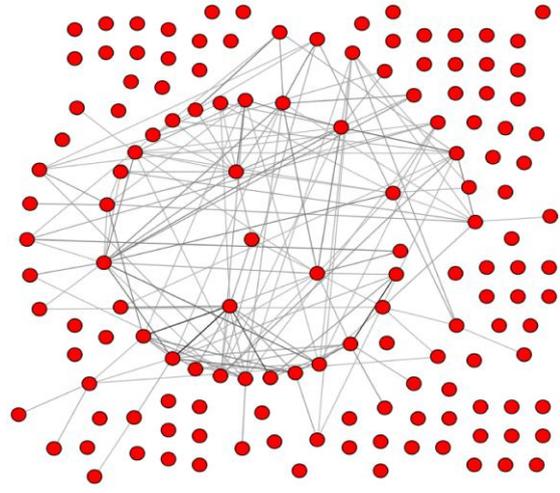
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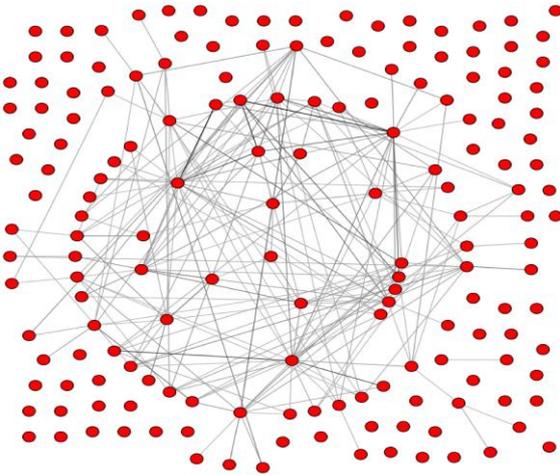
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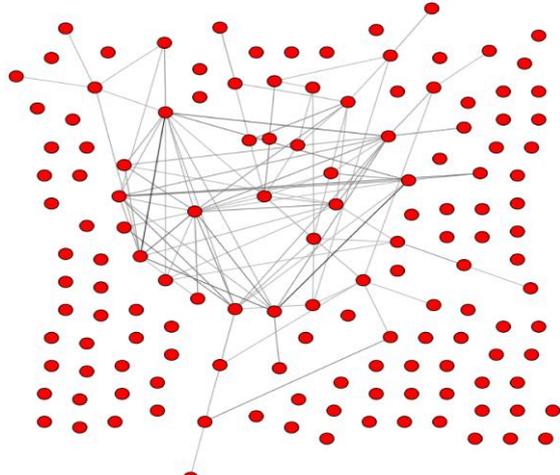
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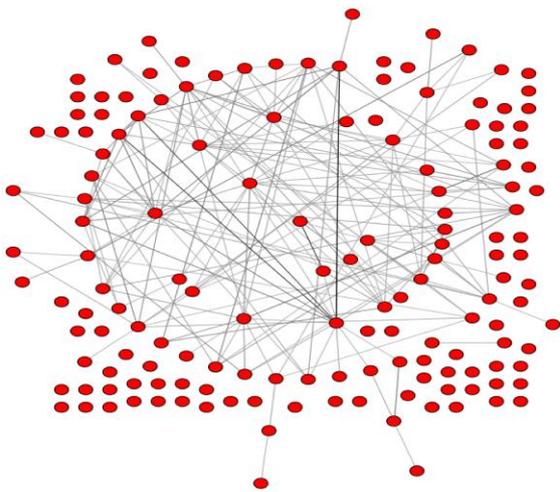
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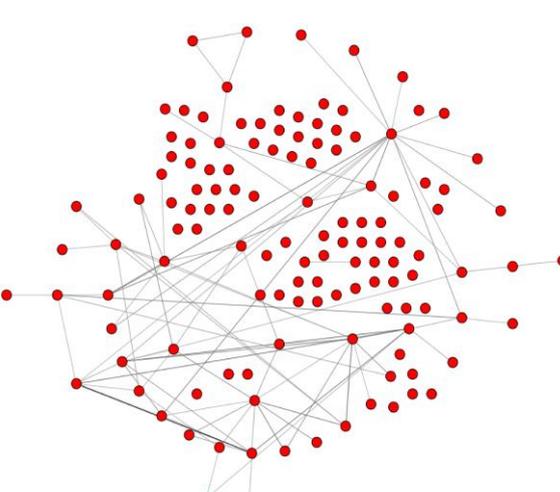
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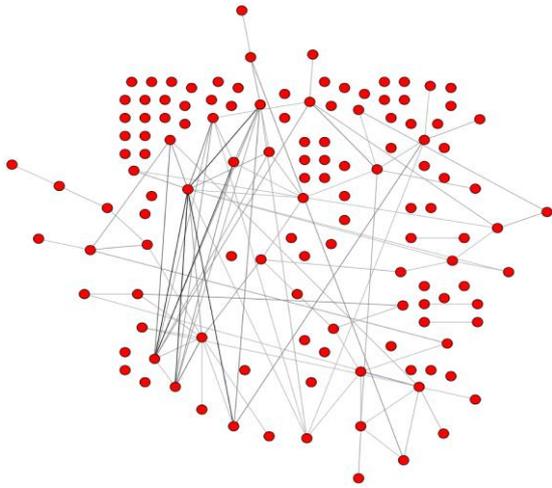
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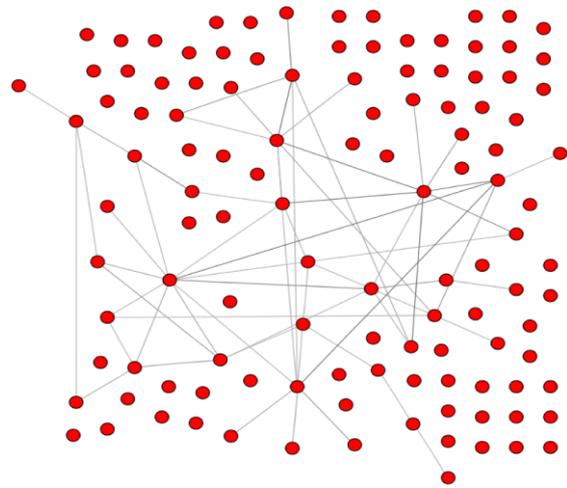
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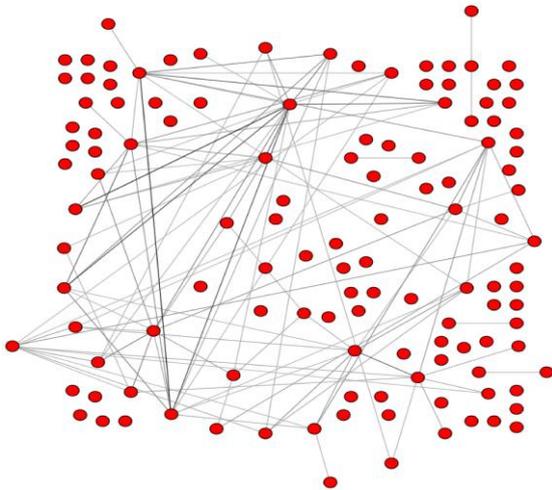
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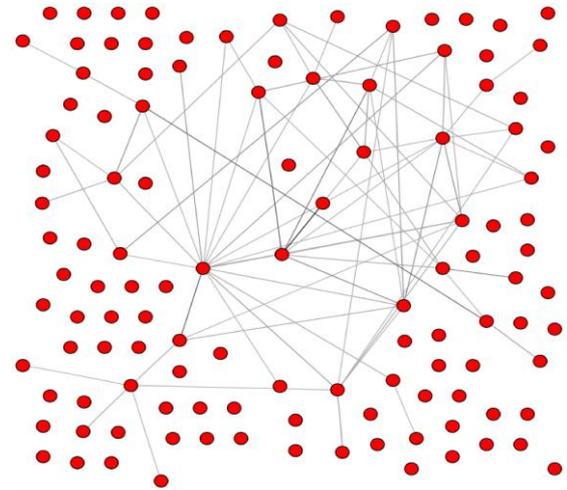
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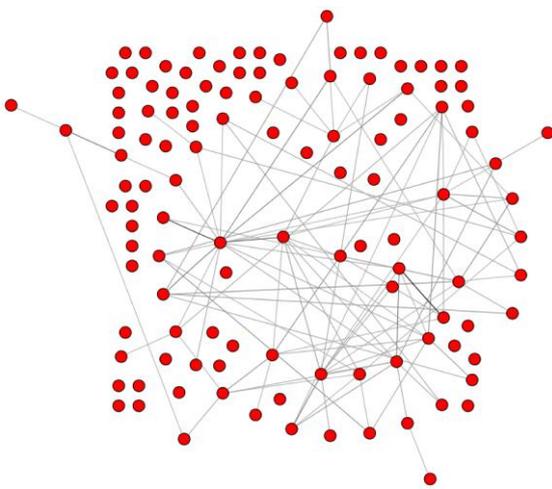
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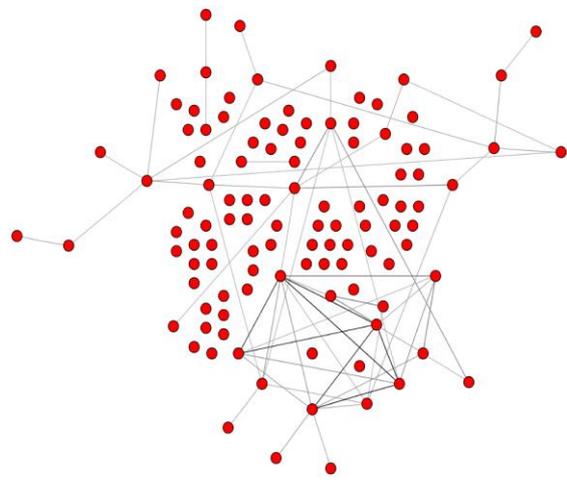
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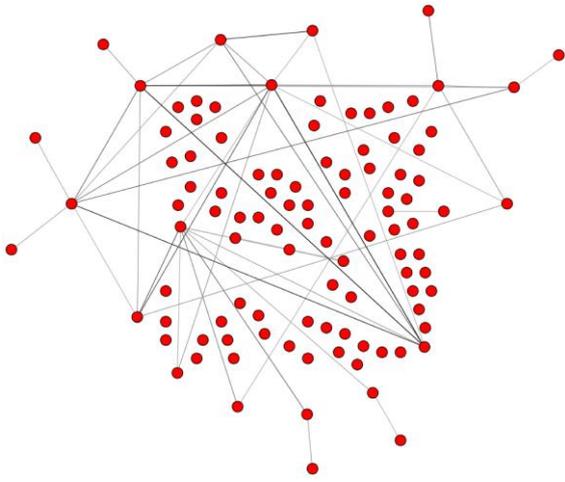
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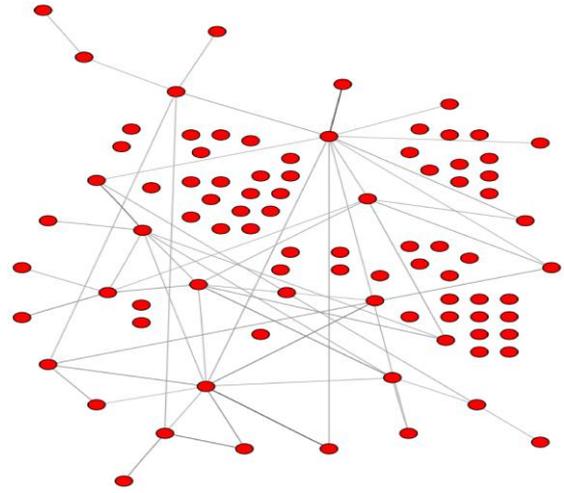
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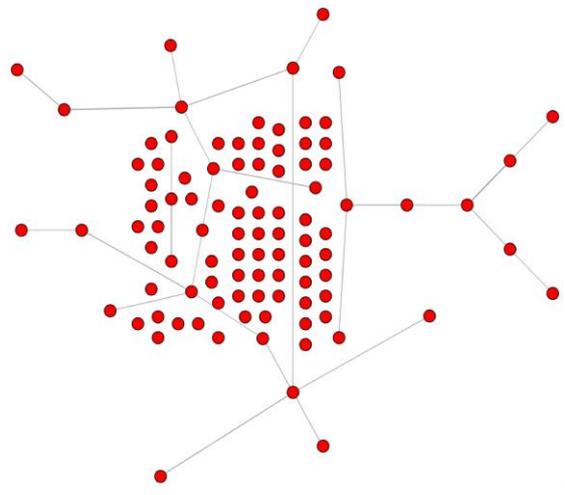
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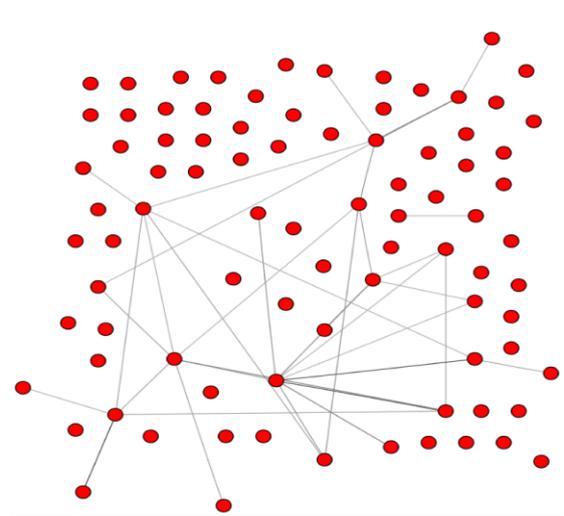
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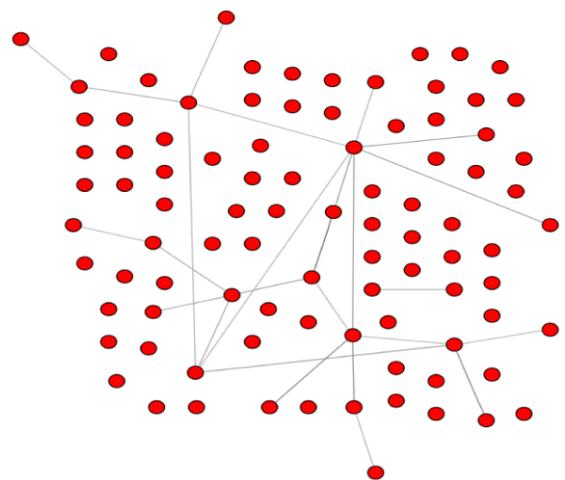
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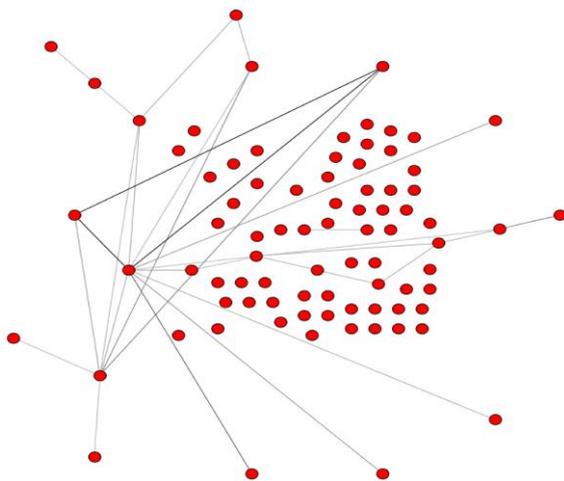
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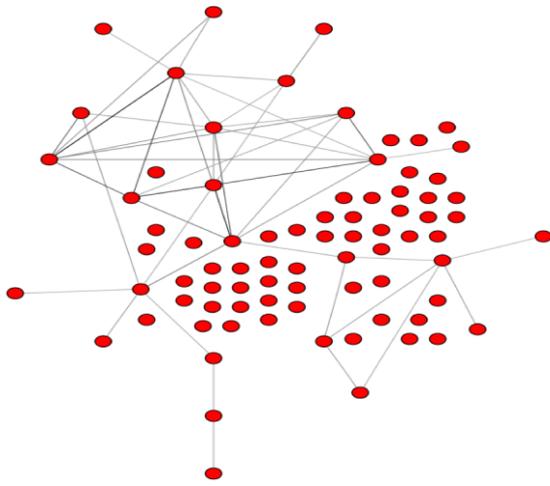
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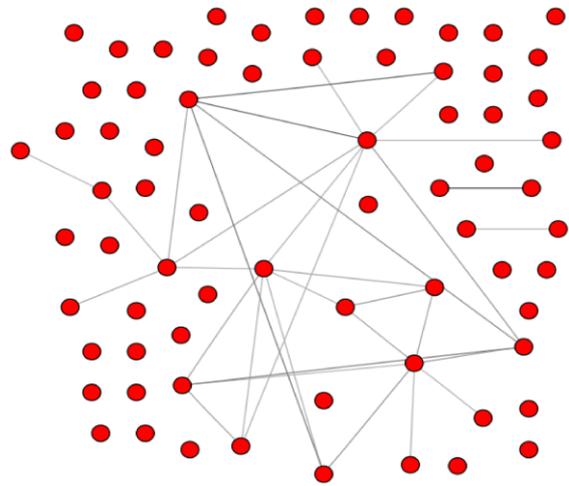
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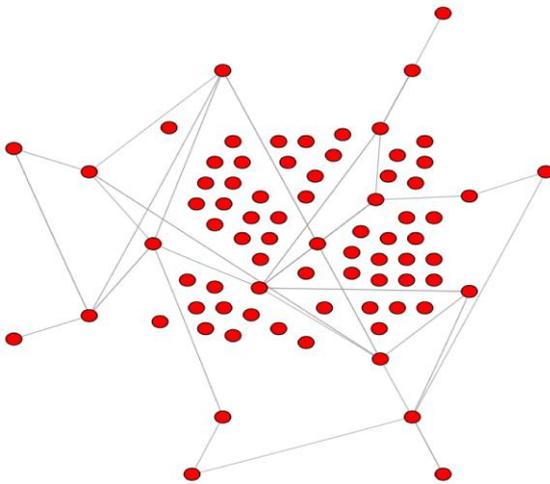
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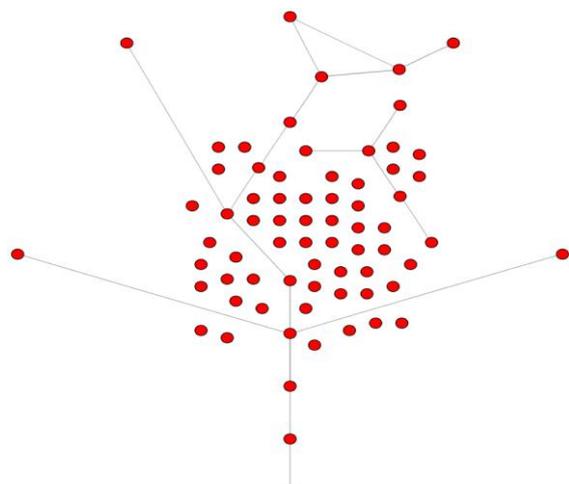
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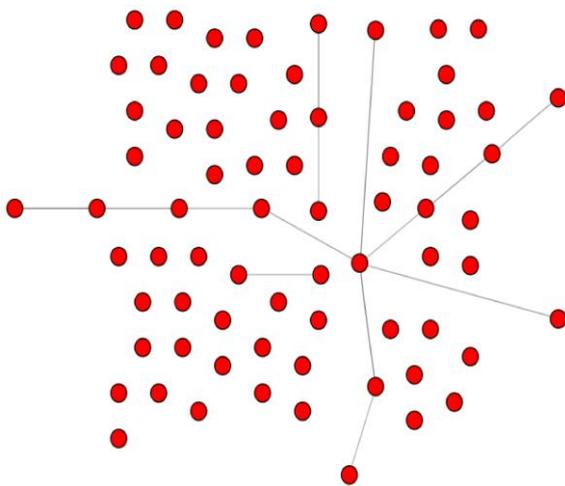
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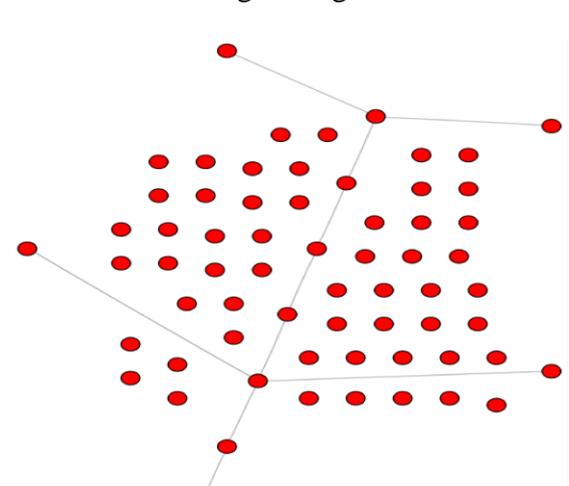
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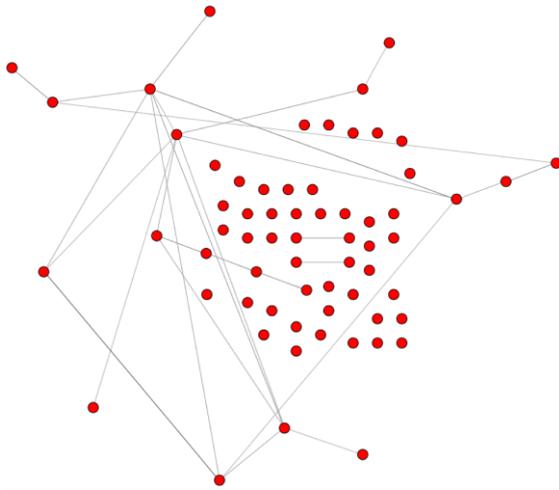
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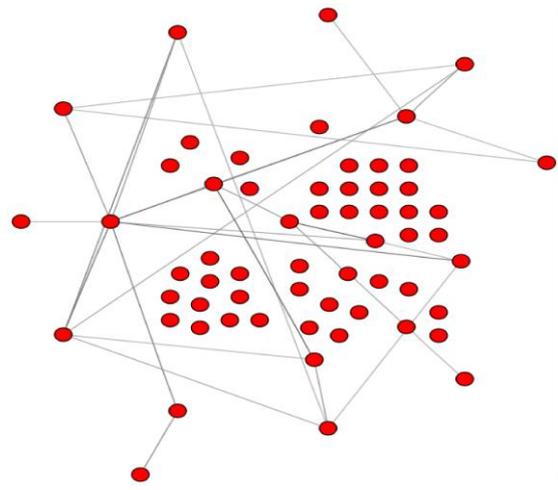
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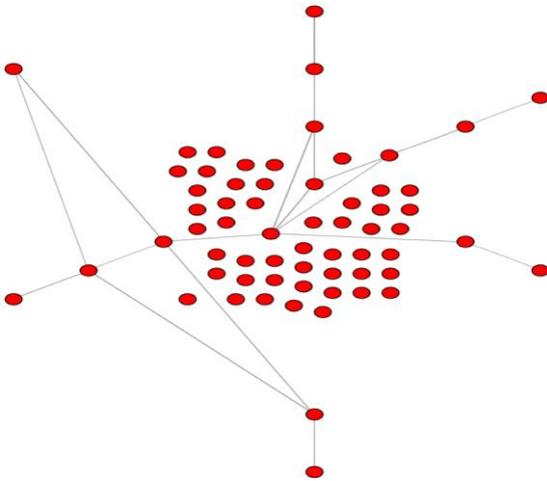
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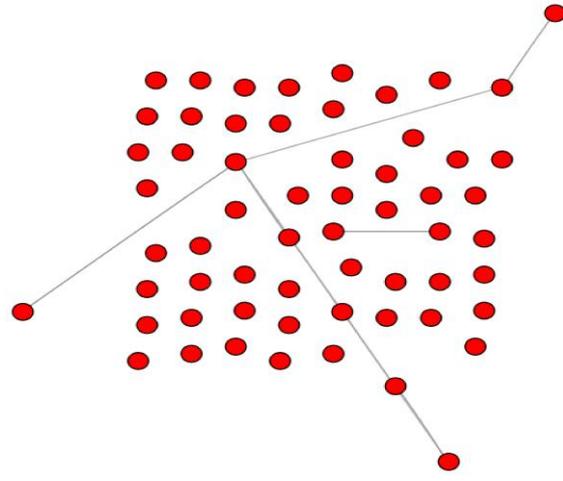
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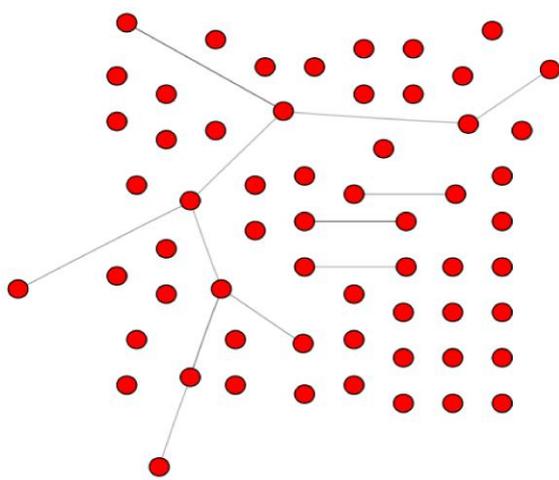
vitamin metabolic process 29 / 60



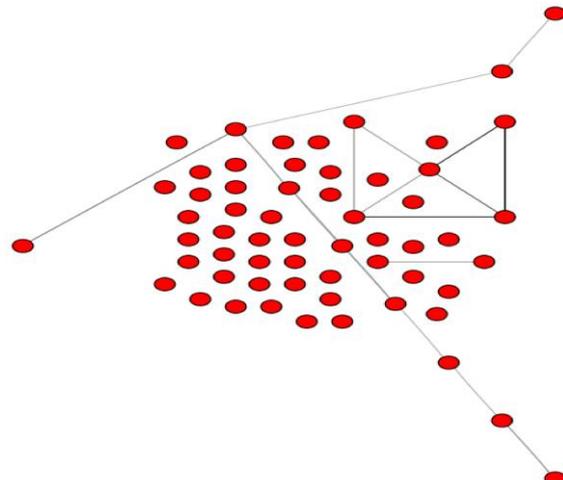
pseudohyphal growth 17 / 63



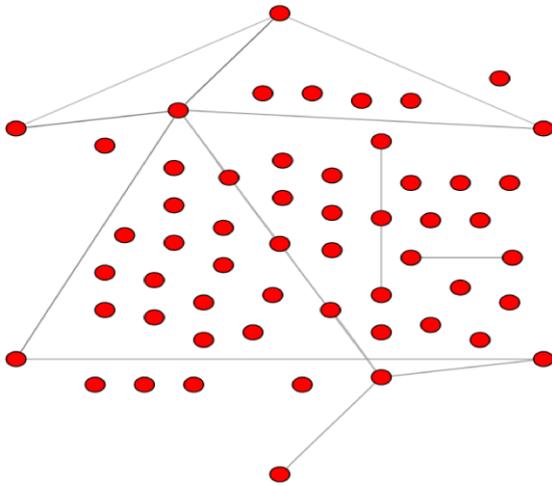
peroxisome 10 / 60



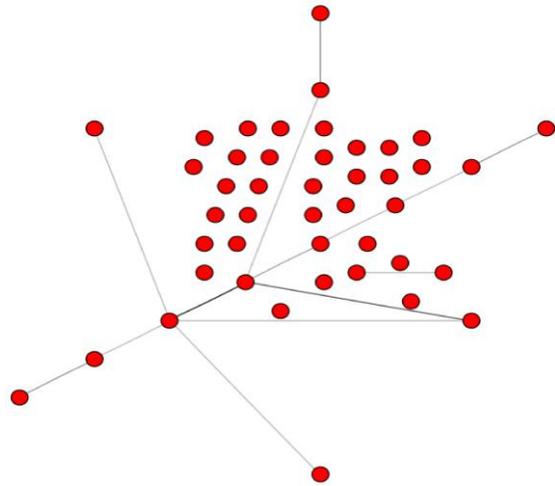
vacuole organization 12 / 60



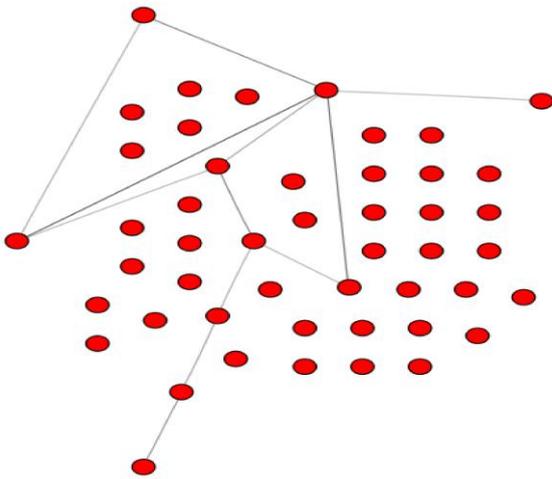
peroxisome organization 18 / 59



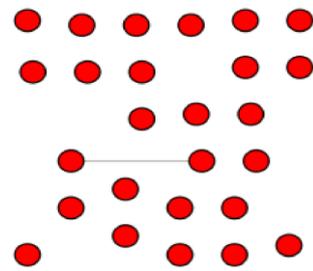
isomerase activity 17 / 55



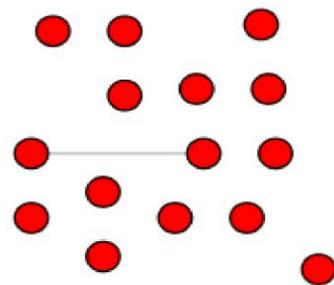
signal transducer activity 14 / 44



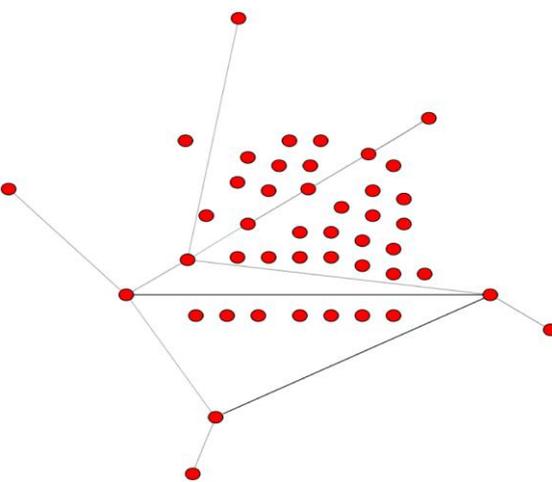
phosphoprotein phosphatase activity 12 / 48



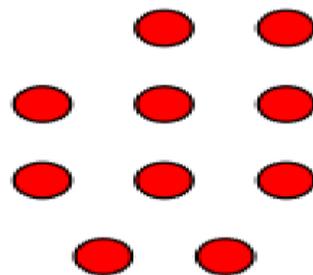
extracellular region 1 / 26



motor activity 1 / 15



transposition 13 / 45



translation regulator activity 0 / 10