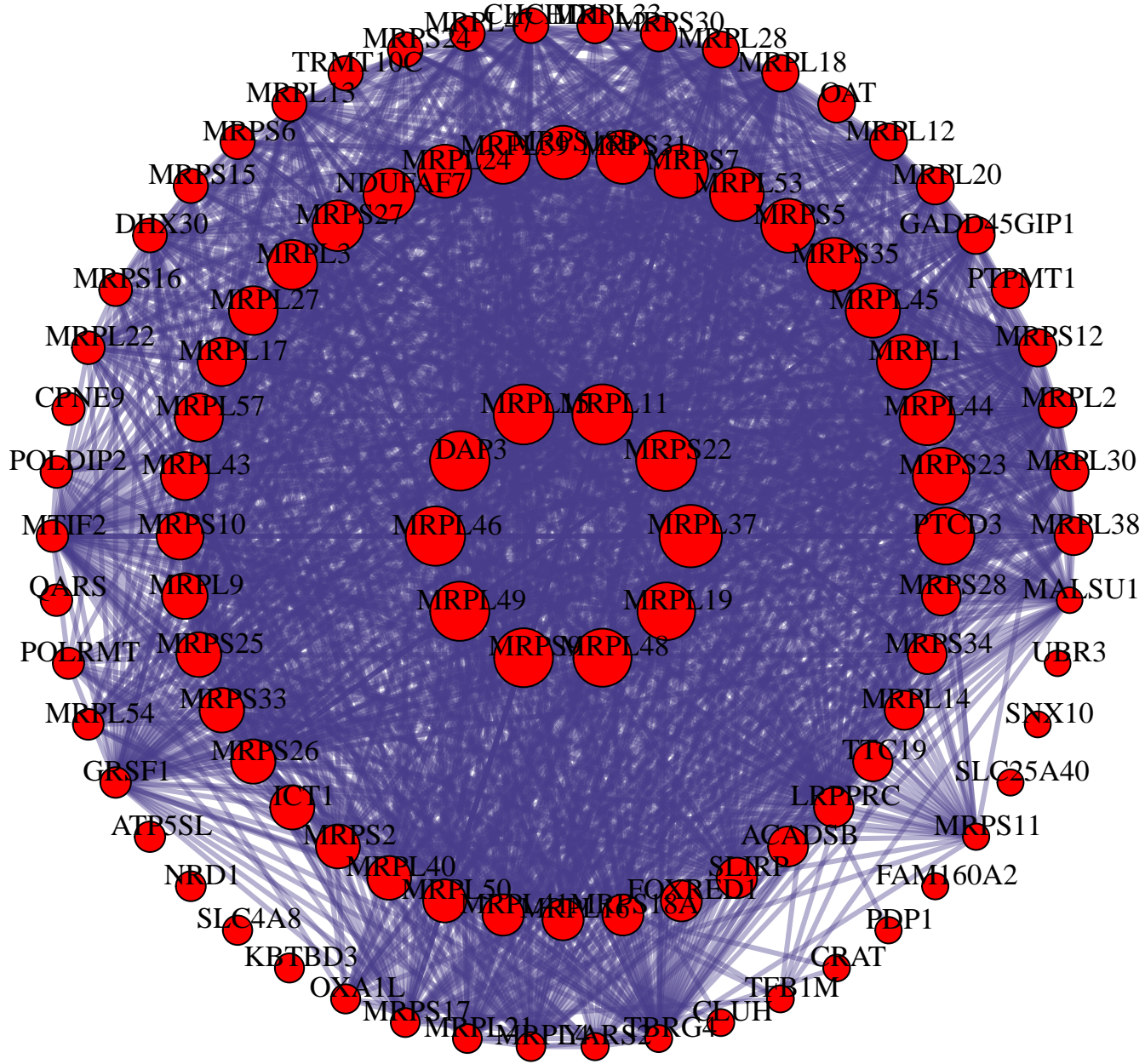
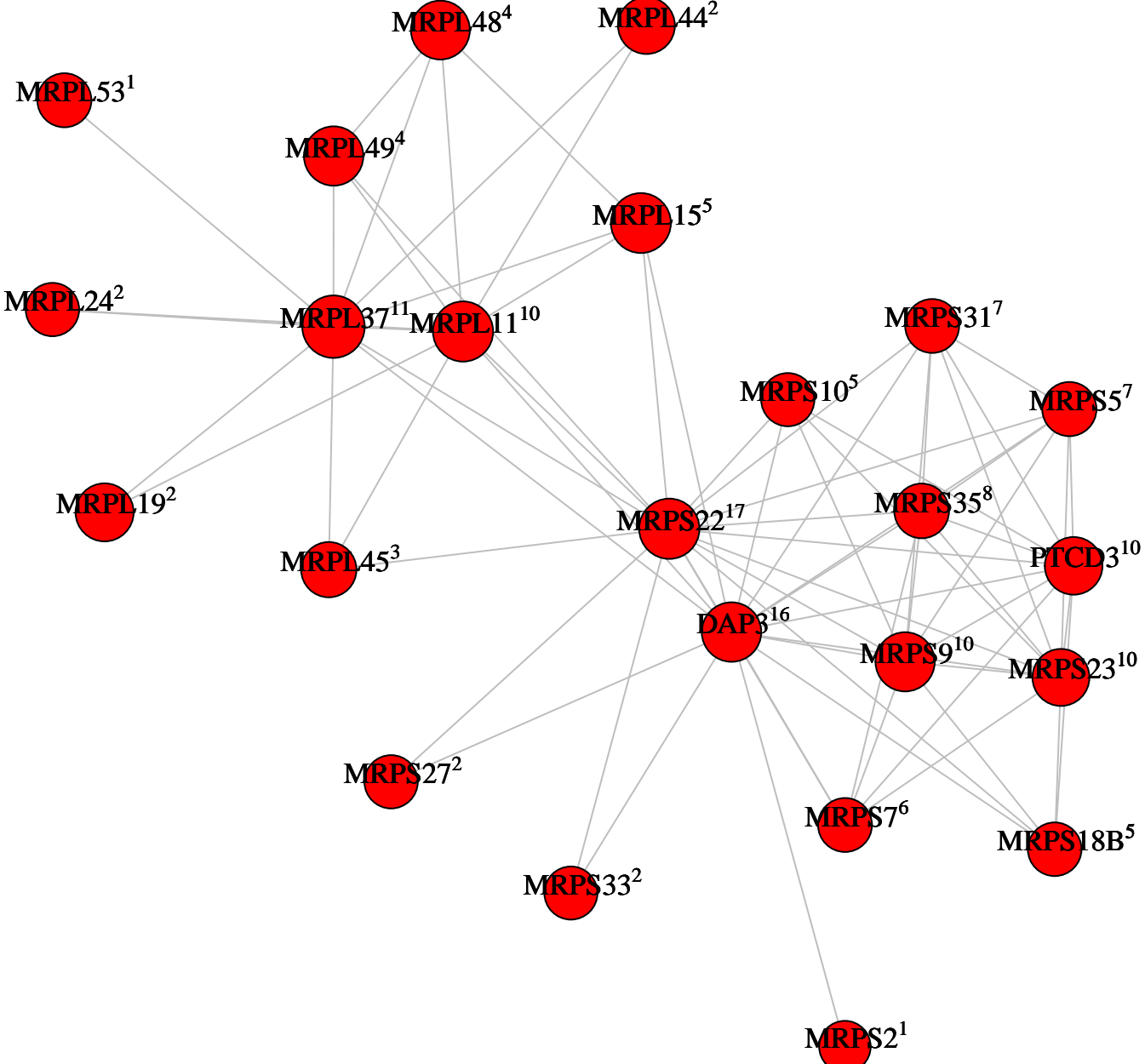


M6 red module



M6 red module hubs connected by top 150 TOM edges: HUB<sup>degree</sup>



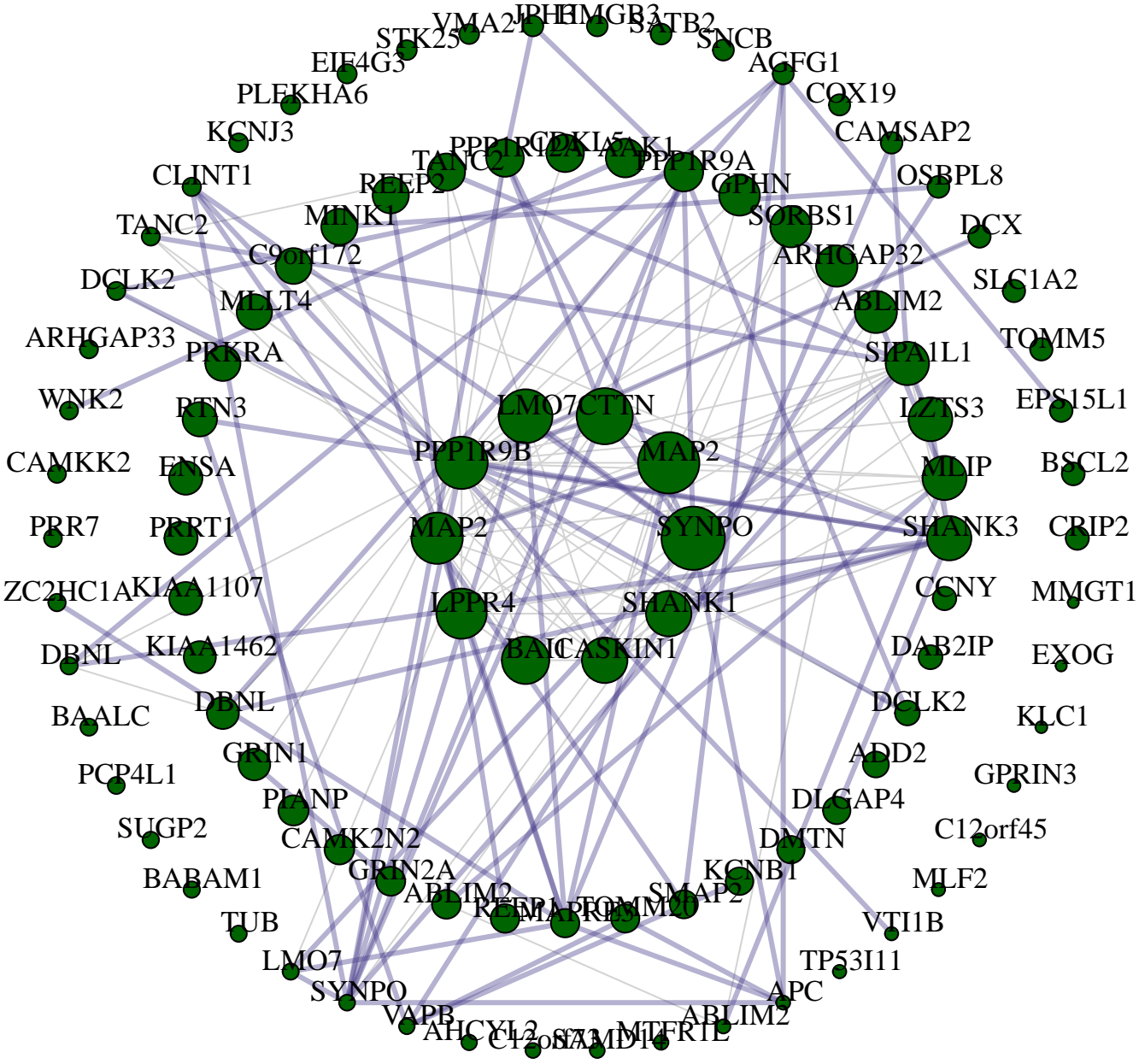
Network diagram showing interactions between various genes. The central node is POLR2B<sup>40</sup>. Other nodes include PRPF6<sup>18</sup>, GCN1L1<sup>10</sup>, POLR2L<sup>3</sup>, POLR2C<sup>4</sup>, TRMT5<sup>6</sup>, UTP14A<sup>2</sup>, CLN5<sup>1</sup>, HBS6<sup>1</sup>, PNKP<sup>1</sup>, ADCK4<sup>1</sup>, CMTR1<sup>1</sup>, MTG1<sup>1</sup>, IFT122<sup>1</sup>, FEN1<sup>1</sup>, USP48<sup>1</sup>, COLEC12<sup>2</sup>, NELFB<sup>1</sup>, SCD<sup>3</sup>, SPPLC1<sup>4</sup>, FKBP10<sup>6</sup>, POLR2J<sup>1</sup>, NQ6<sup>1</sup>, NUDT12<sup>1</sup>, PLXNB3<sup>1</sup>, DENND3<sup>2</sup>, EXOSC10<sup>2</sup>, EMG1<sup>3</sup>, SCP2<sup>2</sup>, FAR1<sup>1</sup>, ZEB2<sup>1</sup>, SCARA3<sup>2</sup>, PCDHGC3<sup>1</sup>, C4orf27<sup>2</sup>, STAT5A<sup>2</sup>, CUL7<sup>1</sup>, NELFA<sup>2</sup>, WDR35<sup>1</sup>, ANKRD39<sup>6</sup>, ARM CX1<sup>4</sup>, and TRIP11<sup>1</sup>.



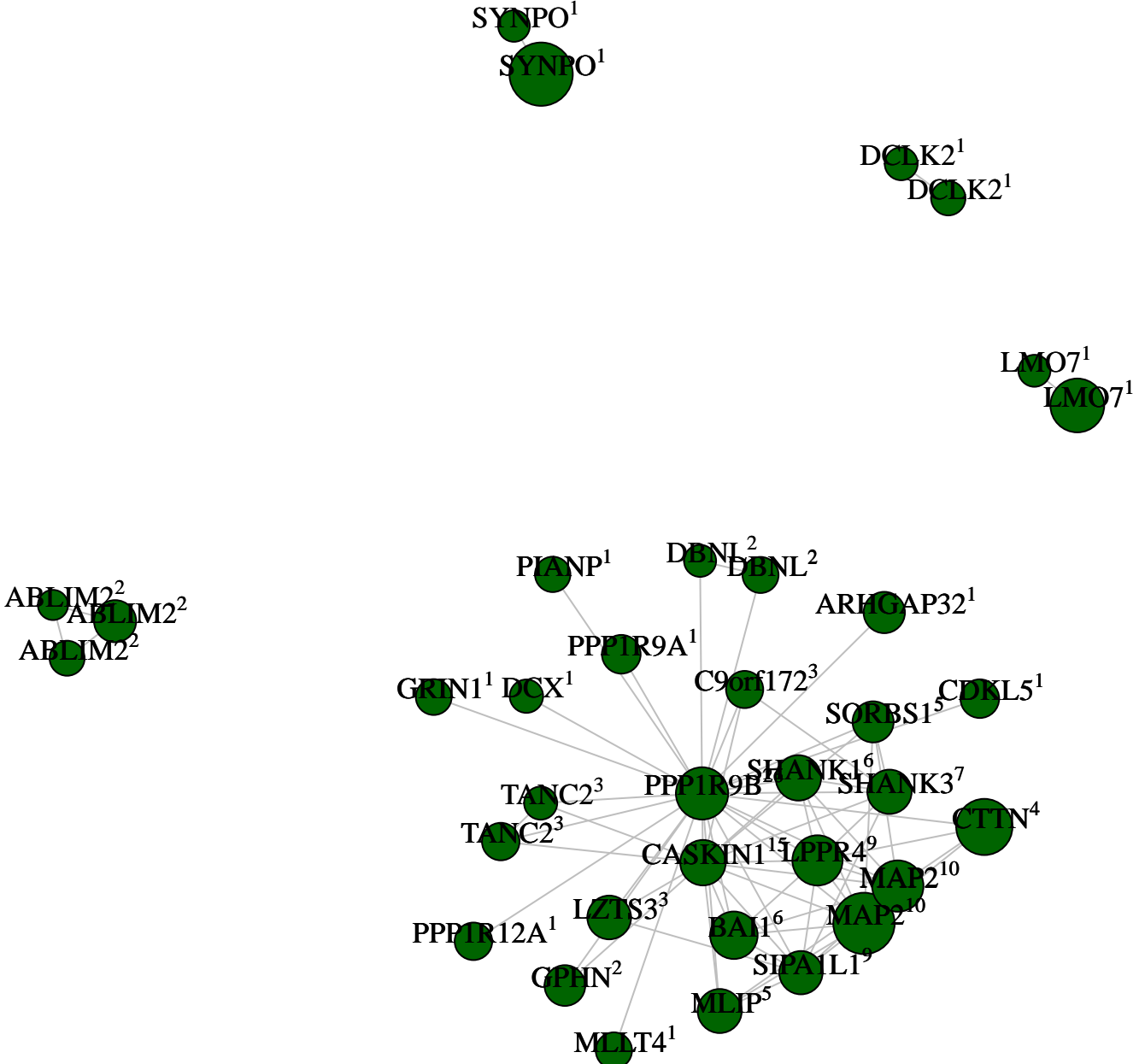
[illegible]



M22 darkgreen module

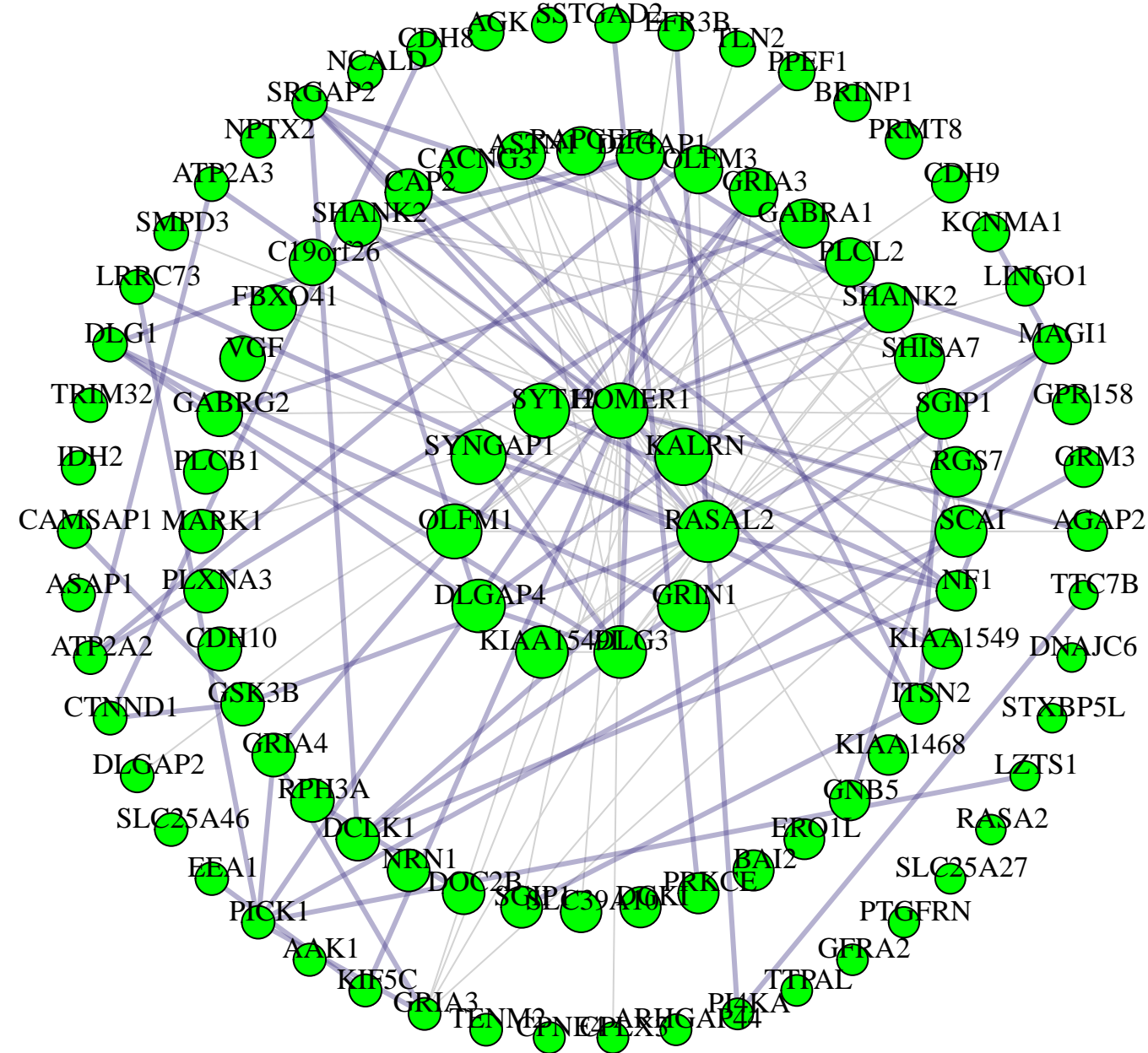


M22 darkgreen module hubs connected by top 150 TOM edges: HUB<sup>degree</sup>

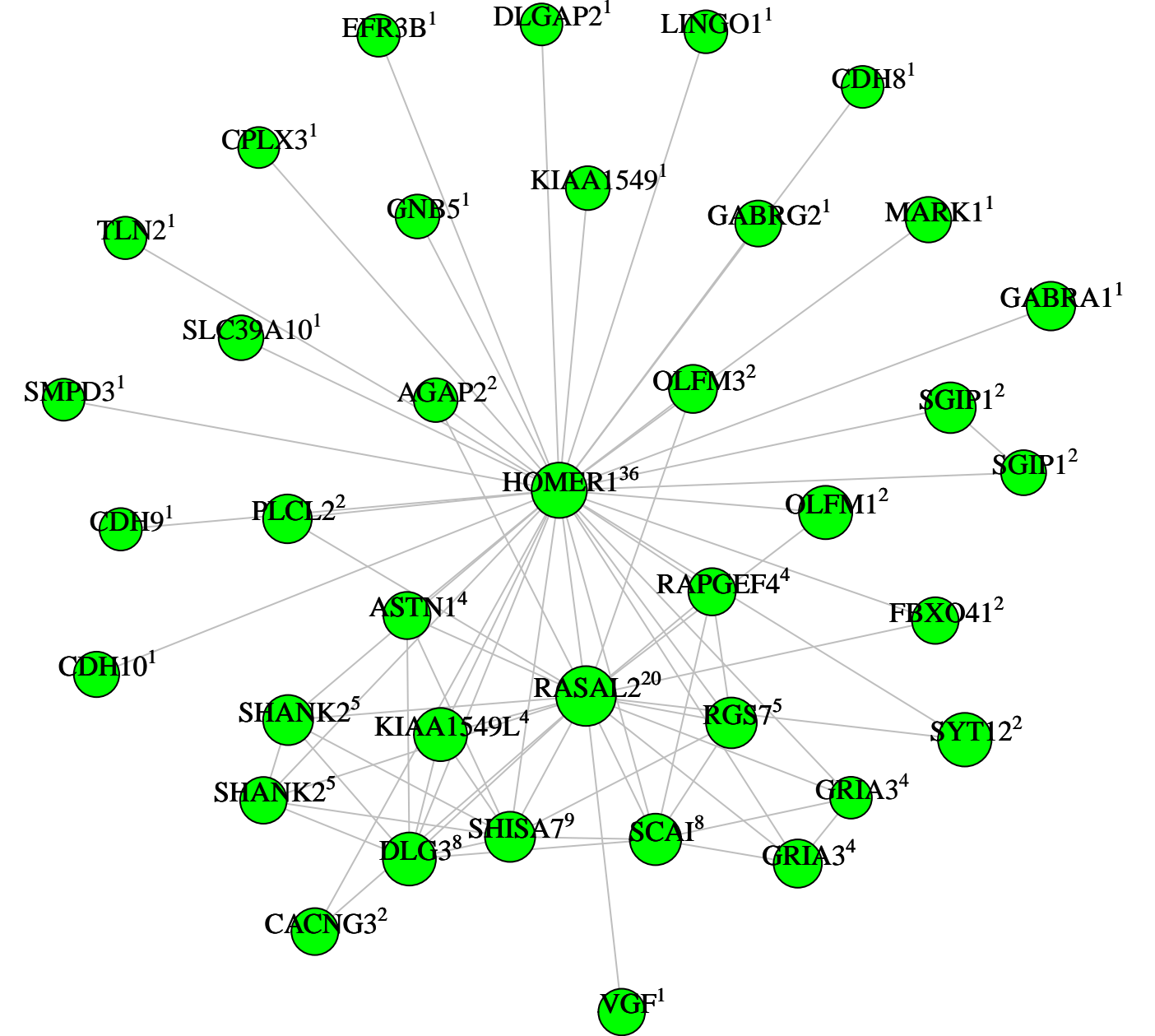




M5 green module



M5 green module hubs connected by top 150 TOM edges: HUB<sup>degree</sup>

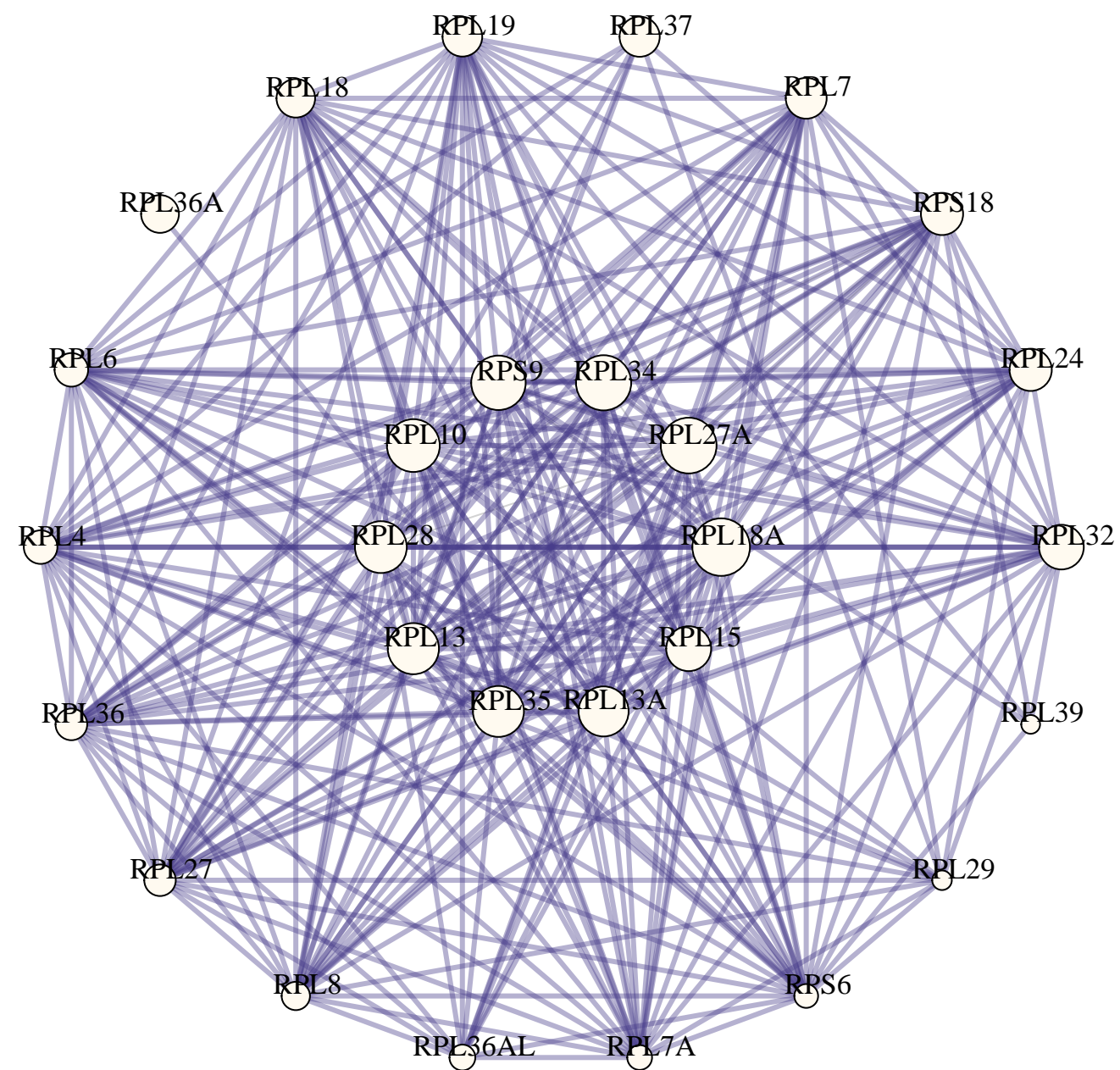


Network diagram showing interactions between various proteins. The nodes are labeled with protein names and superscripted numbers, indicating different isoforms or conditions. The central node is FBXL16, which has the highest degree of connectivity (26). Other highly connected nodes include FBXL16 (26), FBXL16 (26), and FBXL16 (26). The network is dense and interconnected, with many nodes having multiple connections.

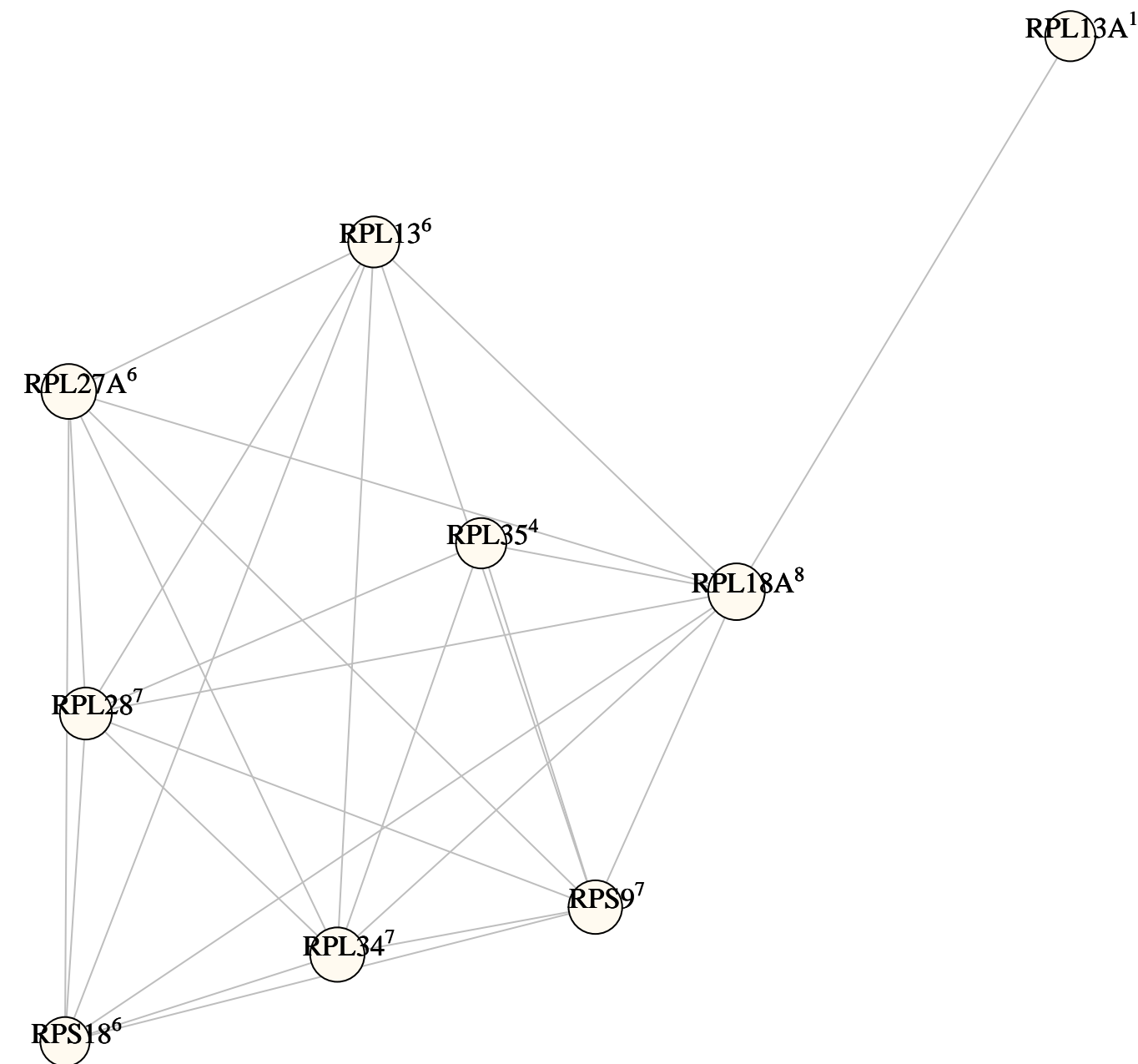
Proteins shown (with their respective counts):

- MPRI1<sup>1</sup>
- MPRI1<sup>1</sup>
- FAM120A<sup>2</sup>
- CAMK4<sup>3</sup>
- ARFGAP2<sup>2</sup>
- USP10<sup>2</sup>
- DOCK3<sup>3</sup>
- CHML<sup>2</sup>
- GAS7<sup>2</sup>
- LARP1<sup>3</sup>
- R3HDM2<sup>6</sup>
- DOCK4<sup>3</sup>
- PPIP5K1<sup>2</sup>
- PPIP5K2<sup>1</sup>
- SNAP25<sup>1</sup>
- KLC2<sup>1</sup>
- FAM73A<sup>1</sup>
- CYB5R1<sup>1</sup>
- SERPINE2<sup>3</sup>
- KIAA1191<sup>3</sup>
- MTMR7<sup>3</sup>
- RASGRF1<sup>14</sup>
- ARHGAP2<sup>2</sup>
- SLC4A7<sup>6</sup>
- RNMT<sup>5</sup>
- OSBPL8<sup>9</sup>
- PGAP1<sup>3</sup>
- ABR4<sup>4</sup>
- DGKH<sup>3</sup>
- NGEF<sup>2</sup>
- LRRC4B<sup>2</sup>
- TRIM9<sup>2</sup>
- FBXL16<sup>26</sup>
- FBXL16<sup>26</sup>
- FBXL16<sup>26</sup>

**M44 floralwhite module**



M44 floralwhite module hubs connected by top 52 TOM edges: HUB<sup>degree</sup>







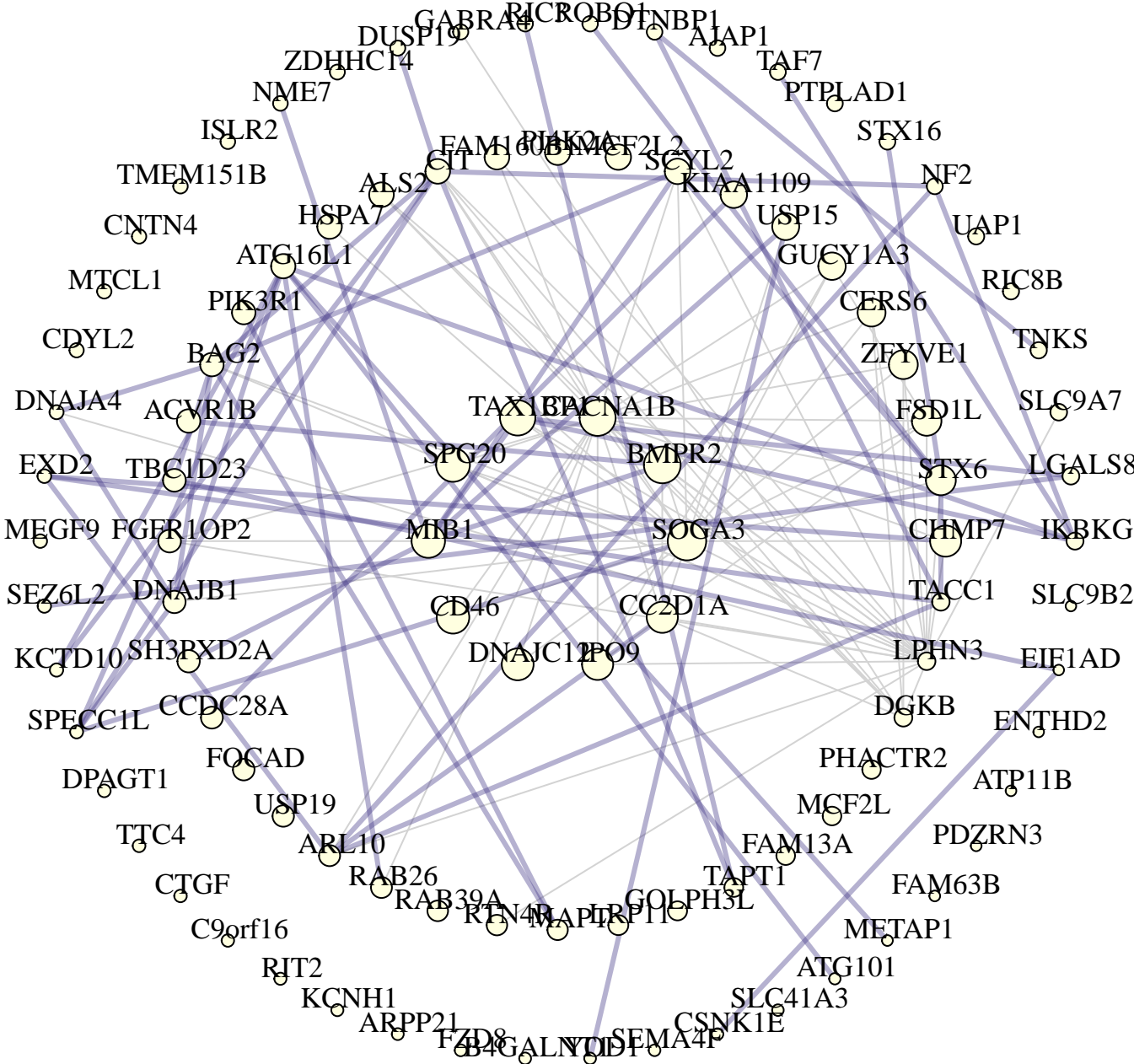




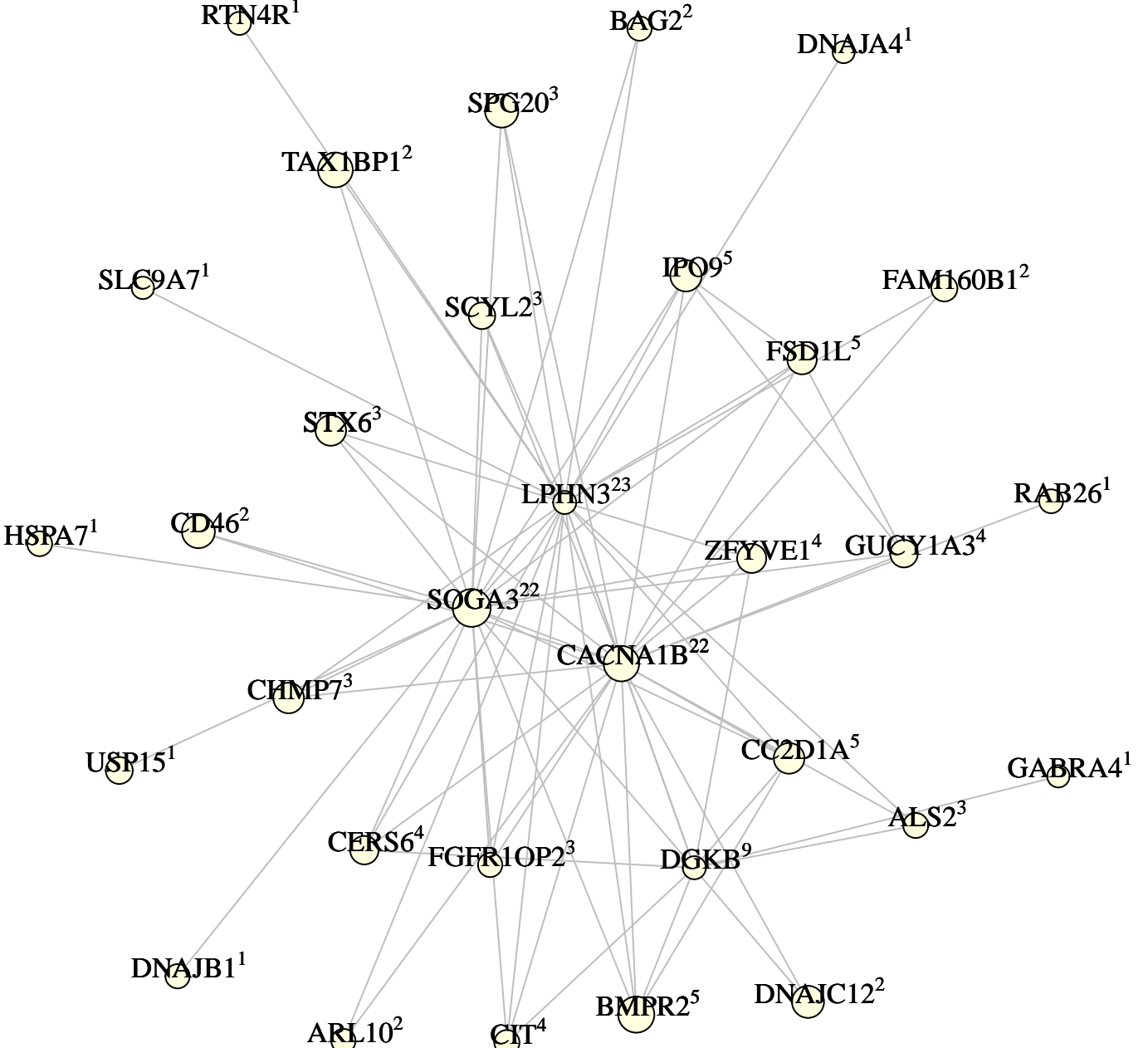


[illegible]

M19 lightyellow module



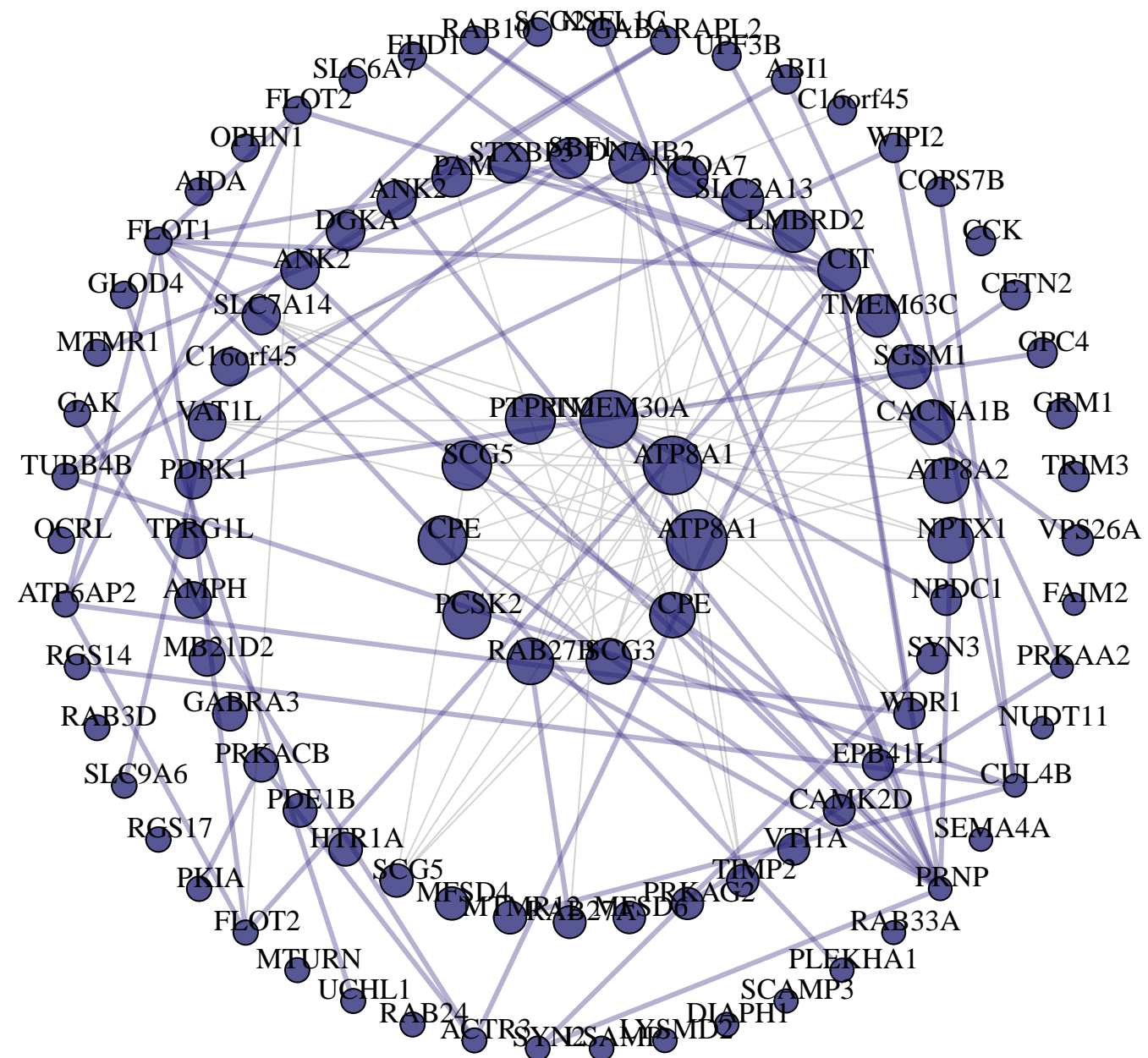
M19 lightyellow module hubs connected by top 150 TOM edges: HUB<sup>degree</sup>



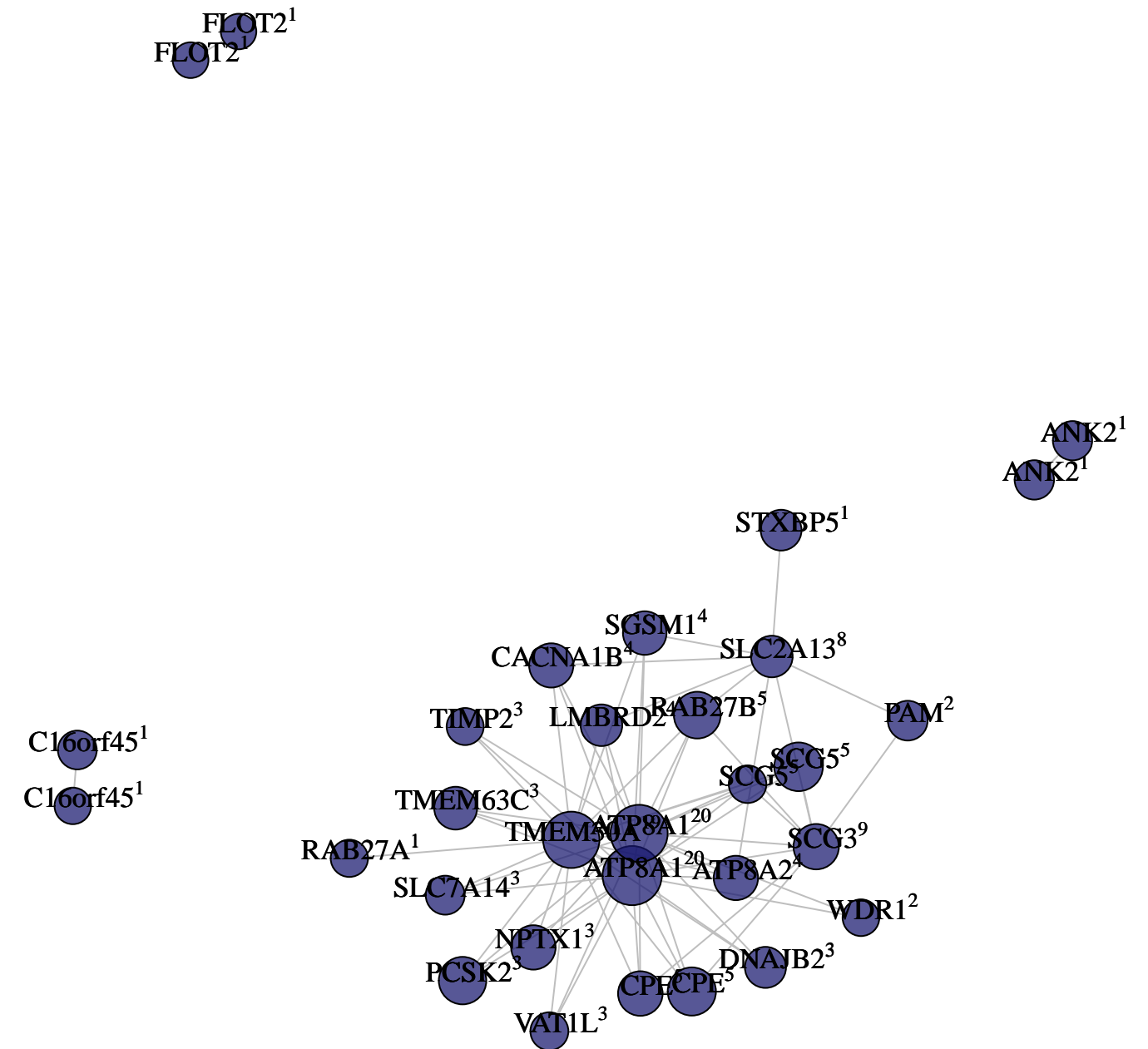




## M15 midnightblue module

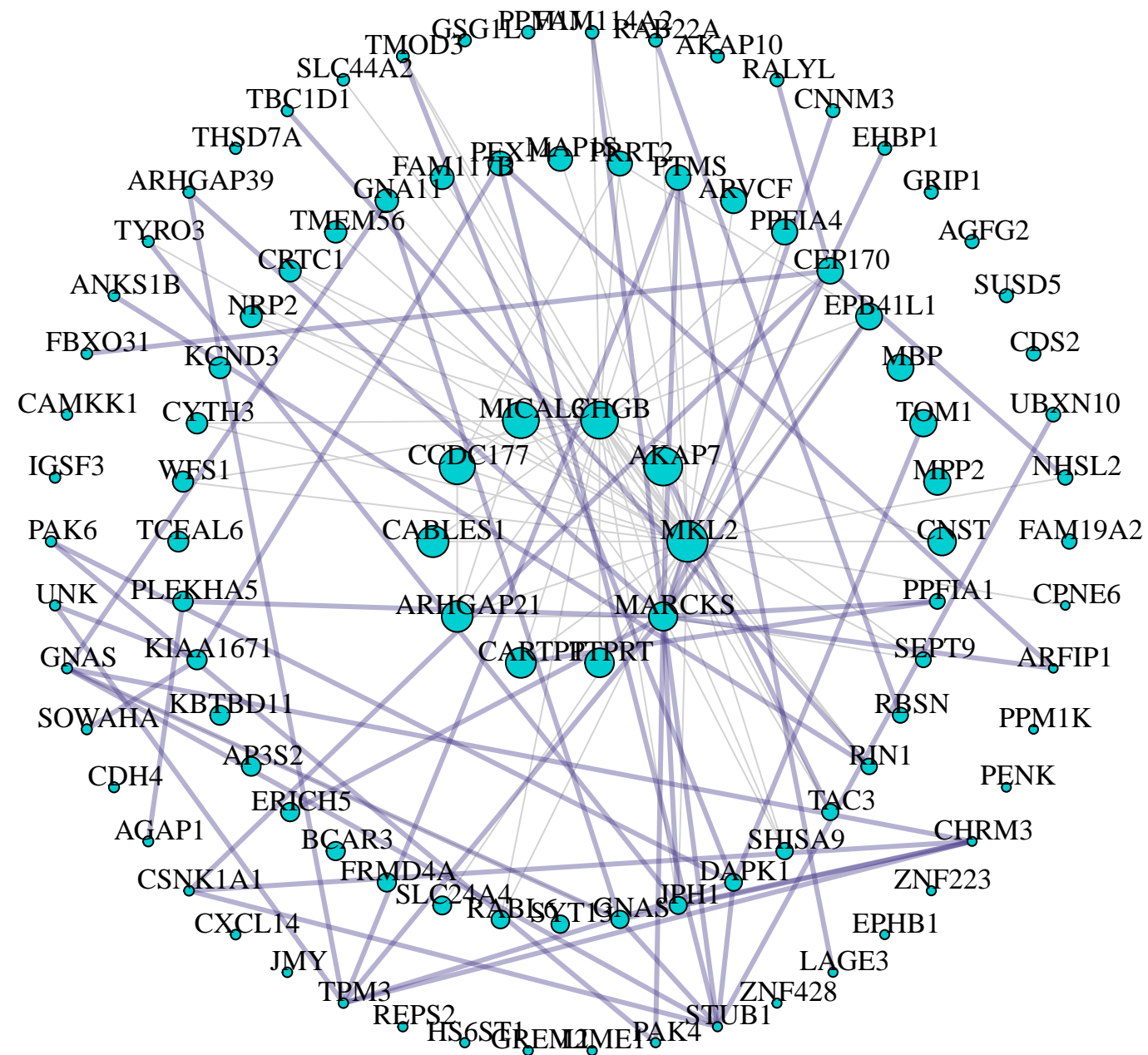


M15 midnightblue module hubs connected by top 150 TOM edges: HUB<sup>degree</sup>

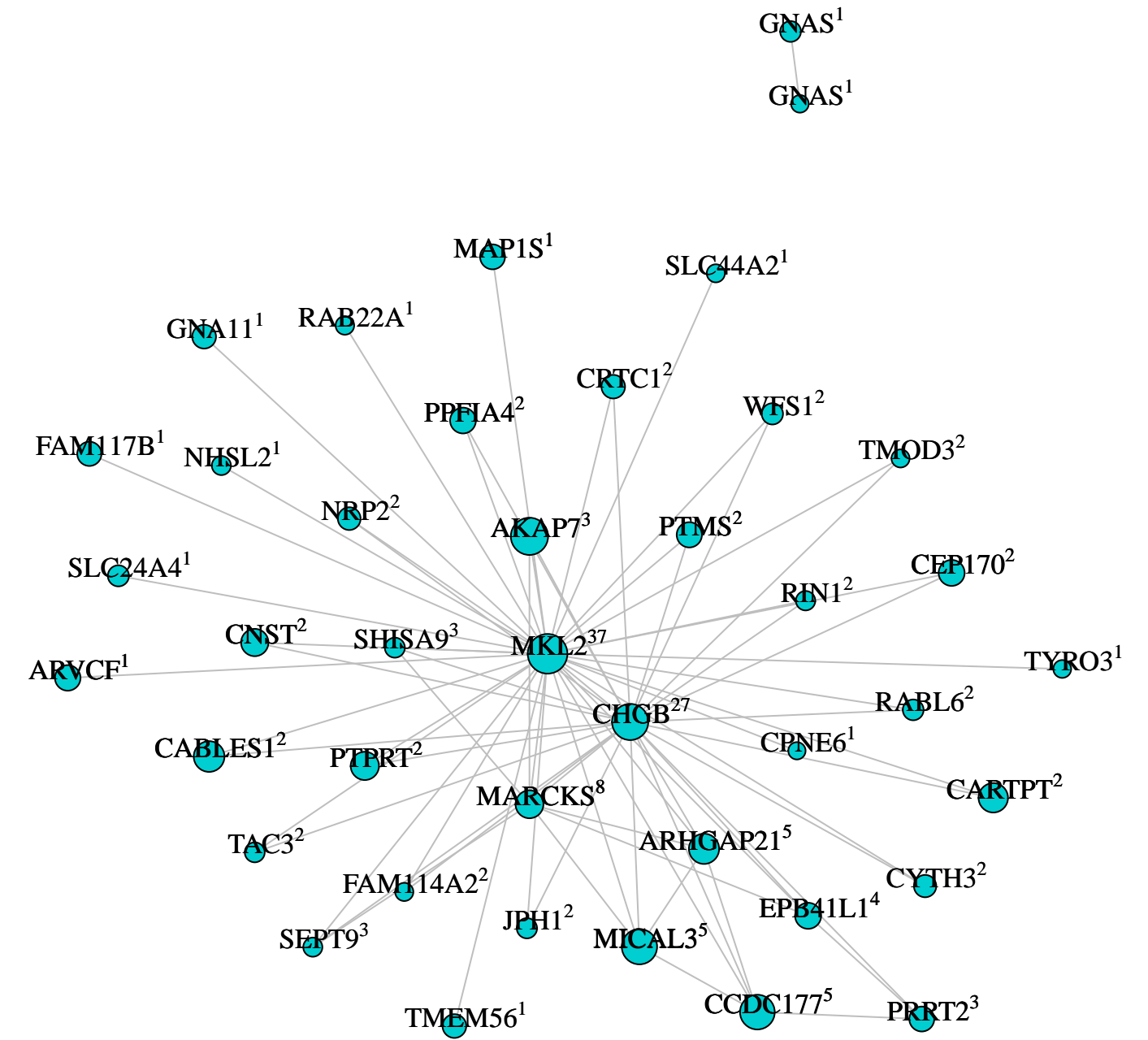


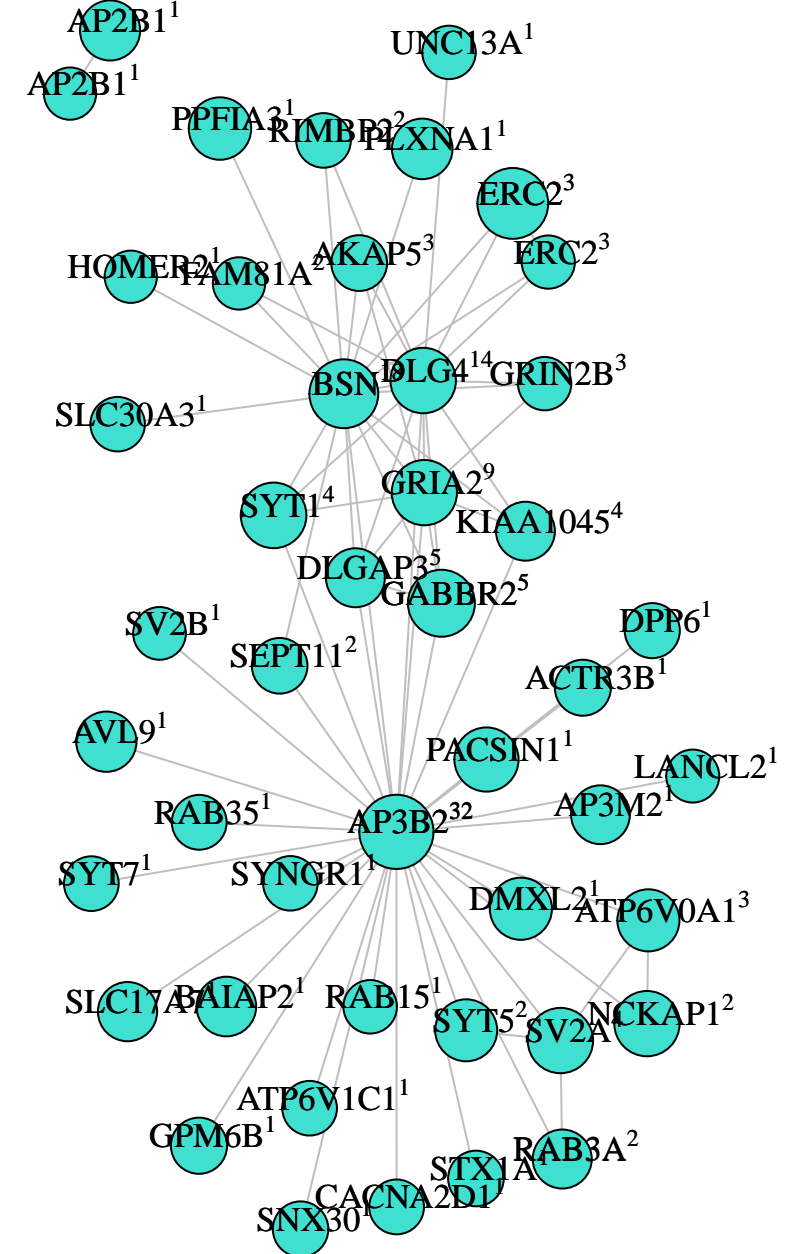


### M23 darkturquoise module



M23 darkturquoise module hubs connected by top 150 TOM edges: HUB<sup>degree</sup>





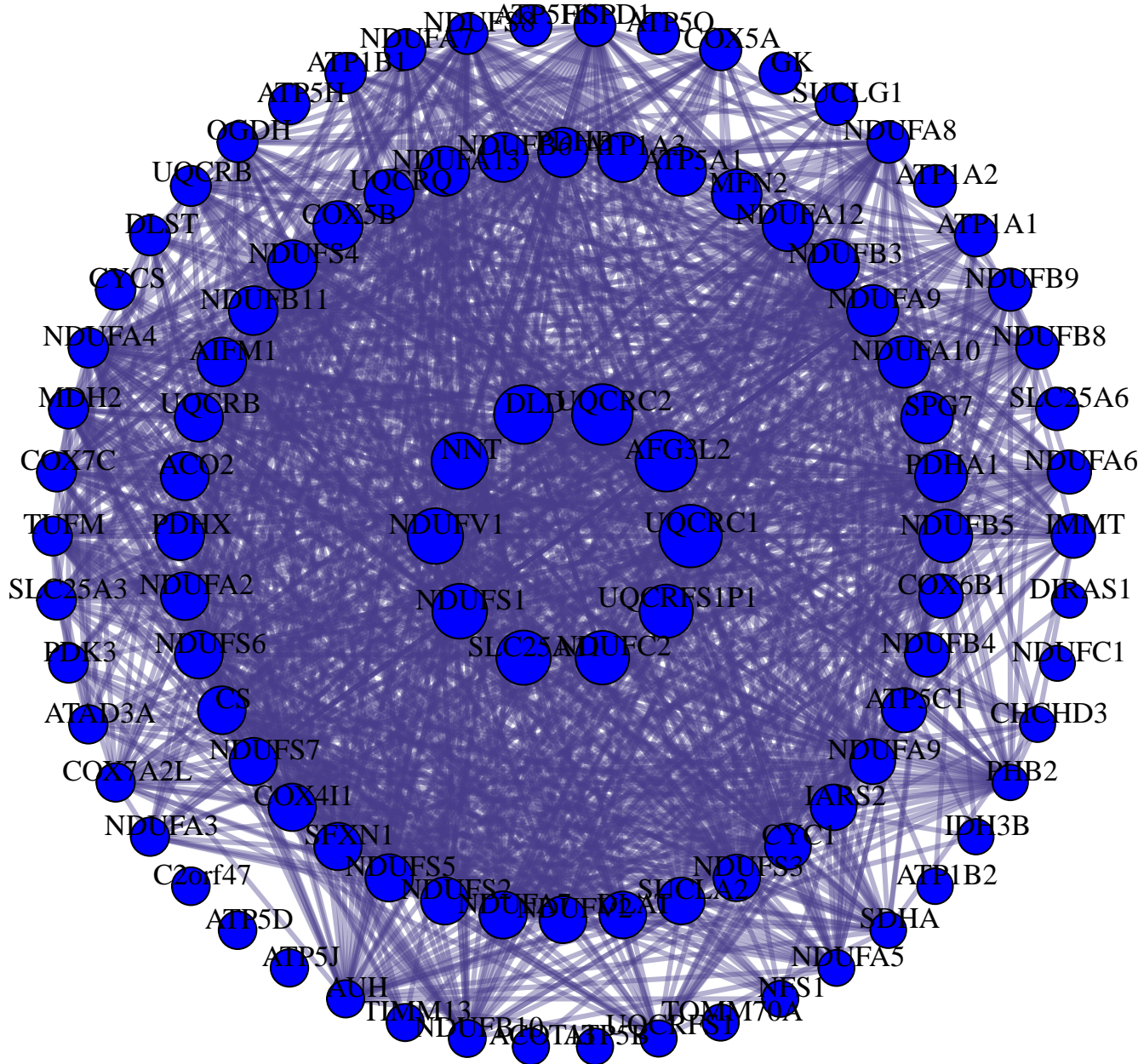


The diagram illustrates a network of protein-protein interactions. The nodes are represented by yellow circles, each labeled with a gene symbol and a superscript indicating the number of interactions or a count. The central node is ATP6V0D1<sup>40</sup>. Other nodes include CLTA<sup>1</sup>, CLTB<sup>1</sup>, AGAP3<sup>1</sup>, CAMK2D<sup>2</sup>, ANK2<sup>1</sup>, and many others. The connections are shown as lines between the nodes, forming a complex web.

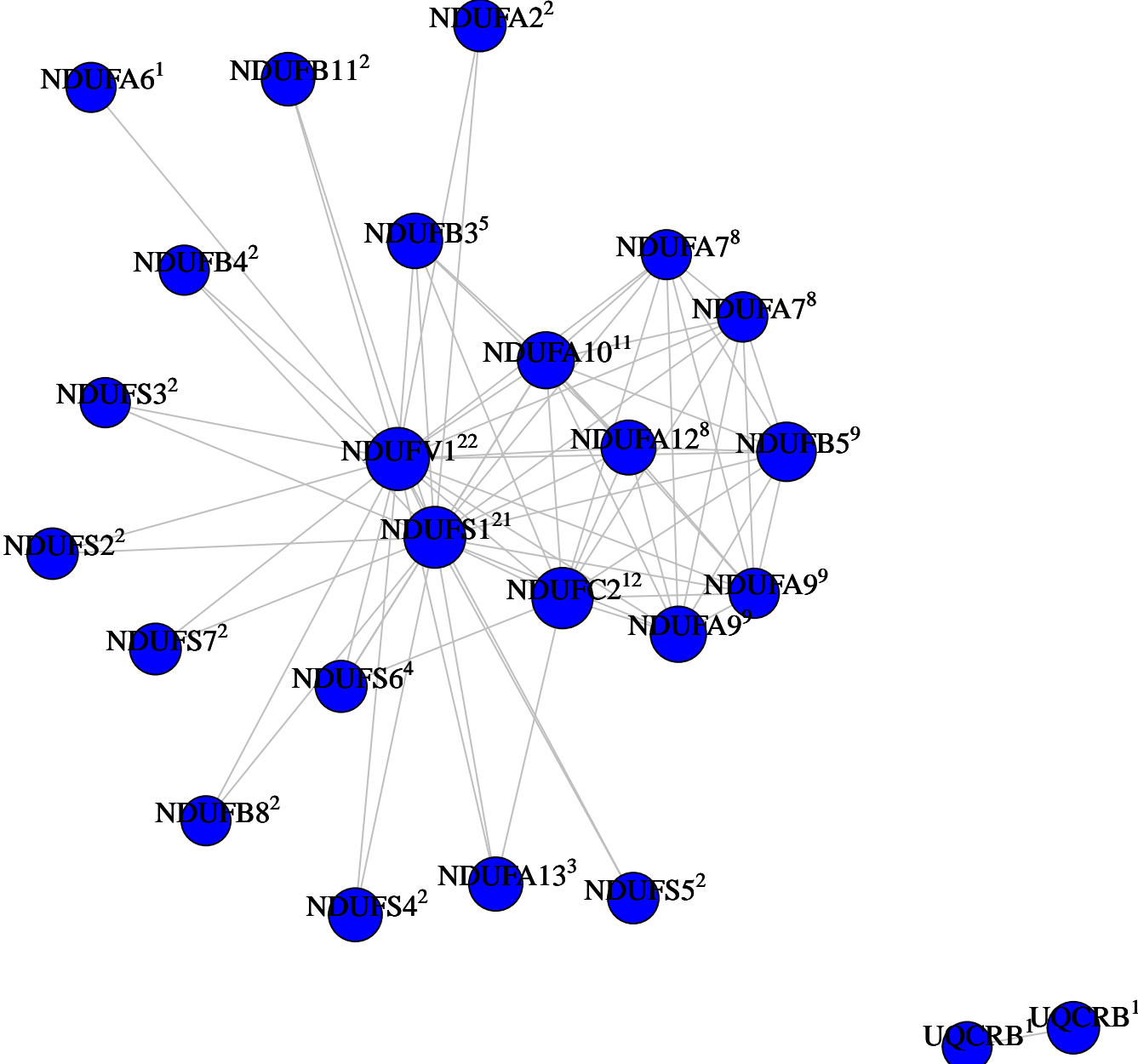
The diagram illustrates a dense network of gene-gene interactions. The nodes, representing genes or proteins, are depicted as green circles. The size of each circle likely corresponds to the degree of connectivity or another quantitative measure. The edges, shown as purple lines, represent the interactions between these entities. The network is characterized by a high density of connections, particularly around central hubs like EGFR, RAB37, SNPH, CAMK2G, and SLC6A1. The overall structure suggests a complex biological pathway or a highly integrated signaling network.



M2 blue module

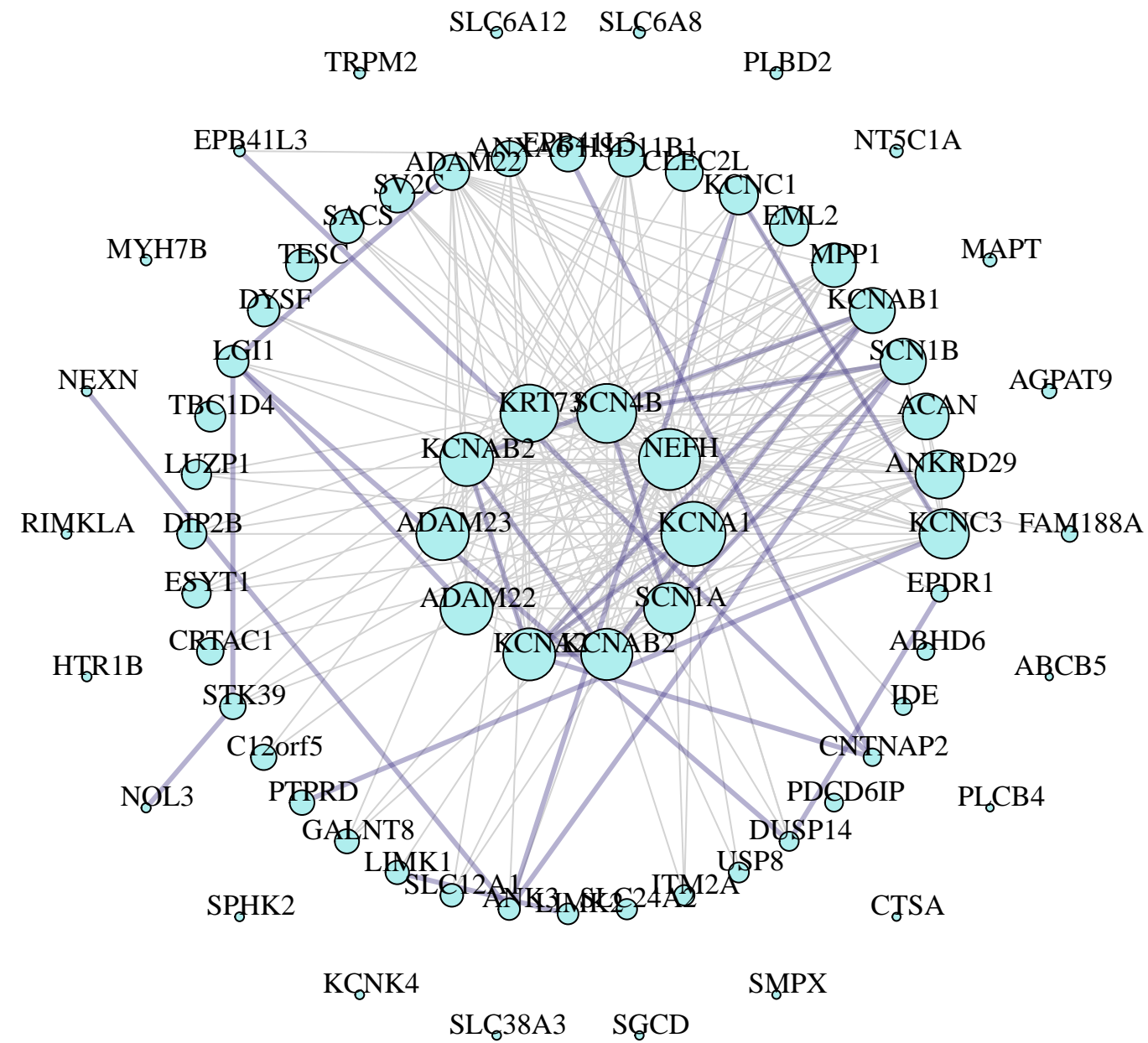


M2 blue module hubs connected by top 150 TOM edges: HUB<sup>degree</sup>

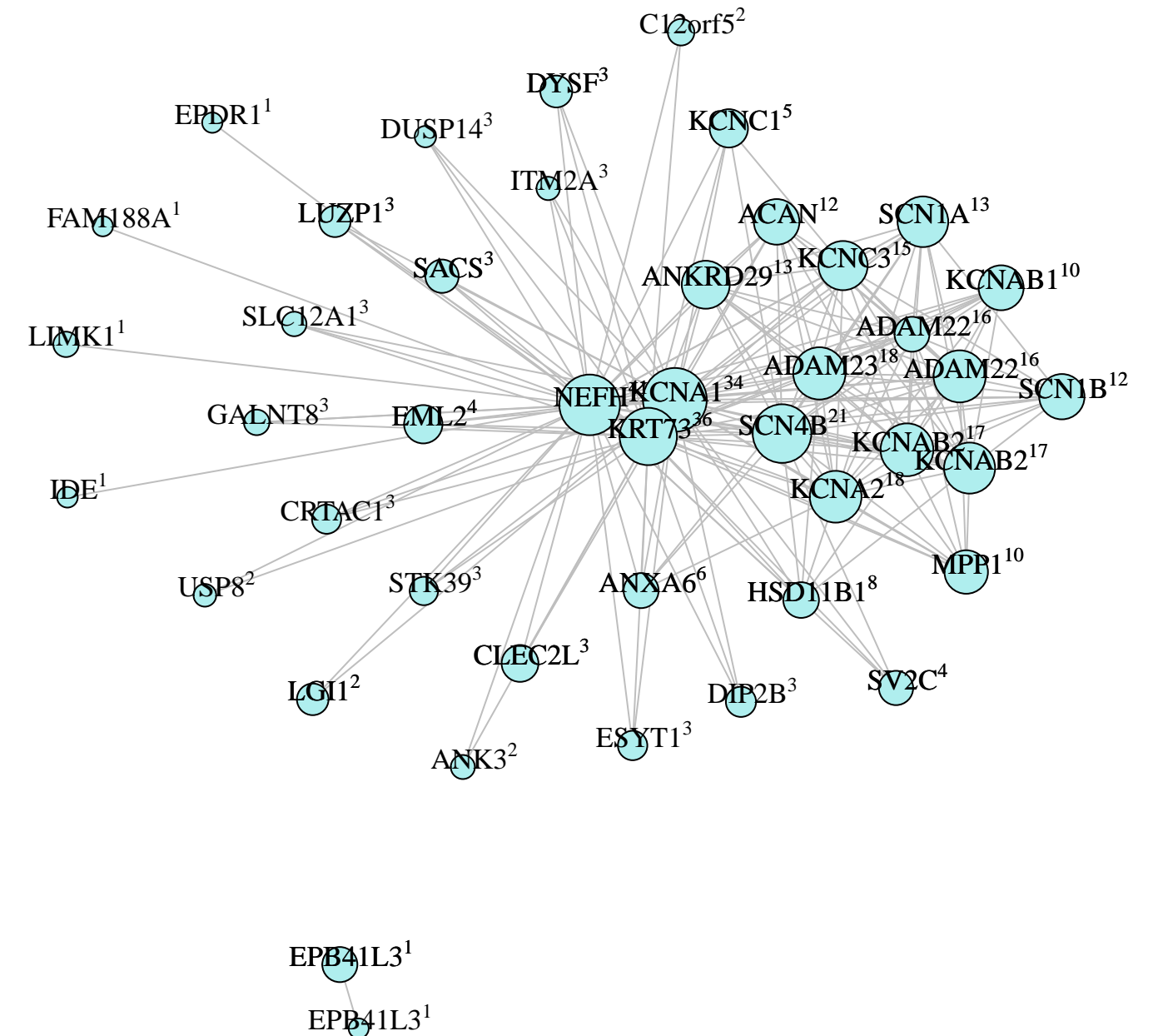




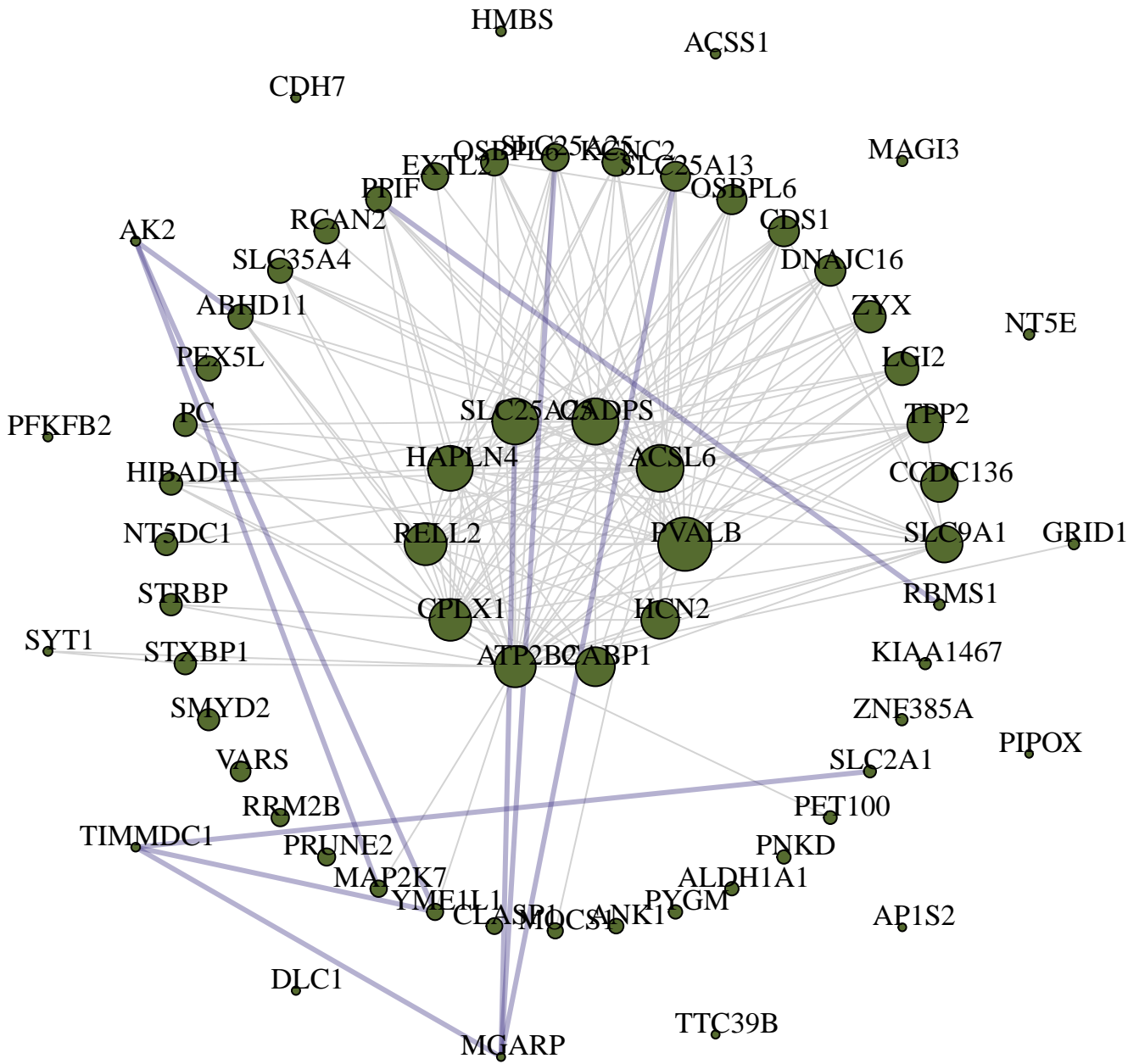
### M31 paleturquoise module



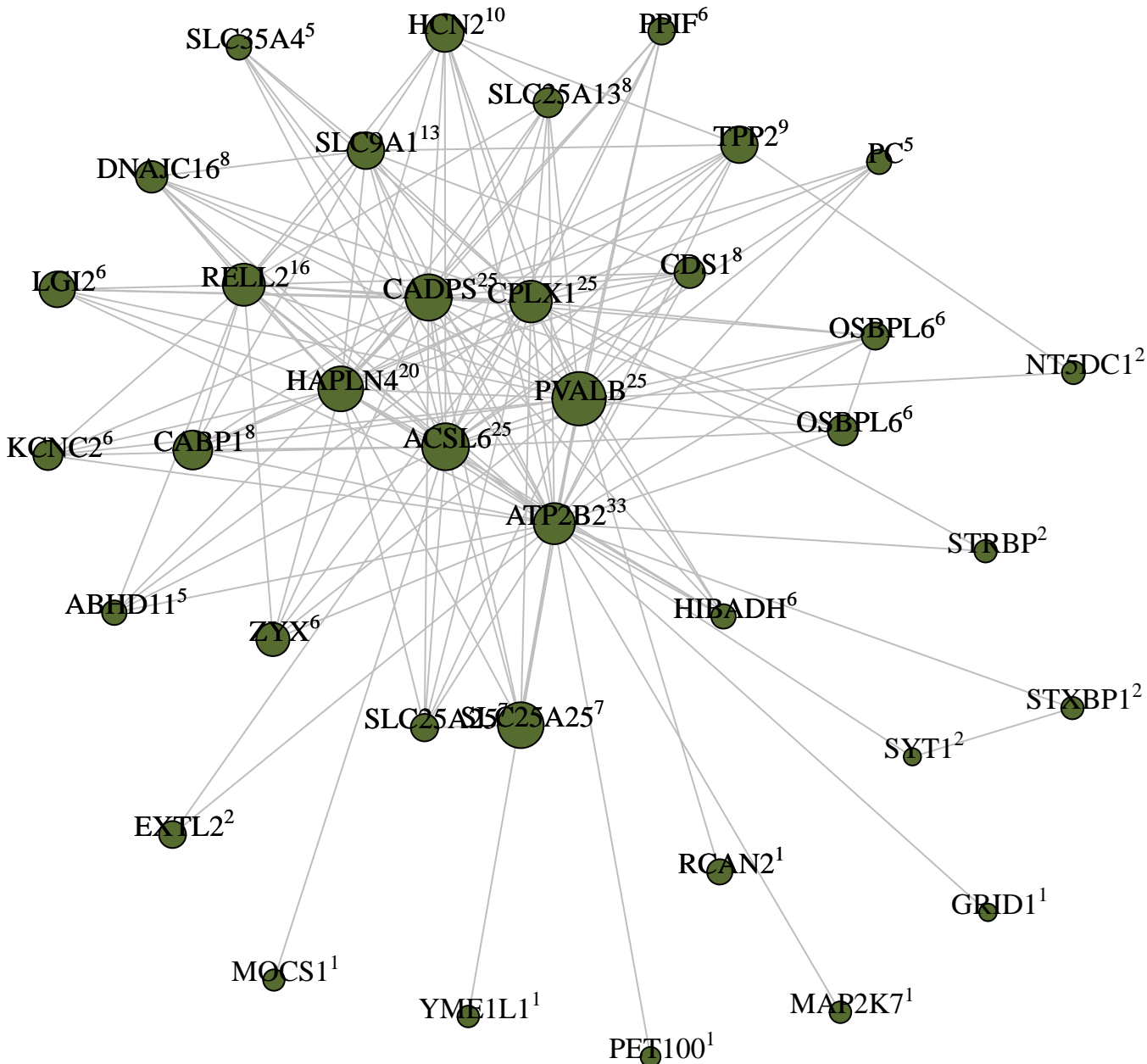
M31 paleturquoise module hubs connected by top 396 TOM edges: HUB<sup>degree</sup>



M33 darkolivegreen module

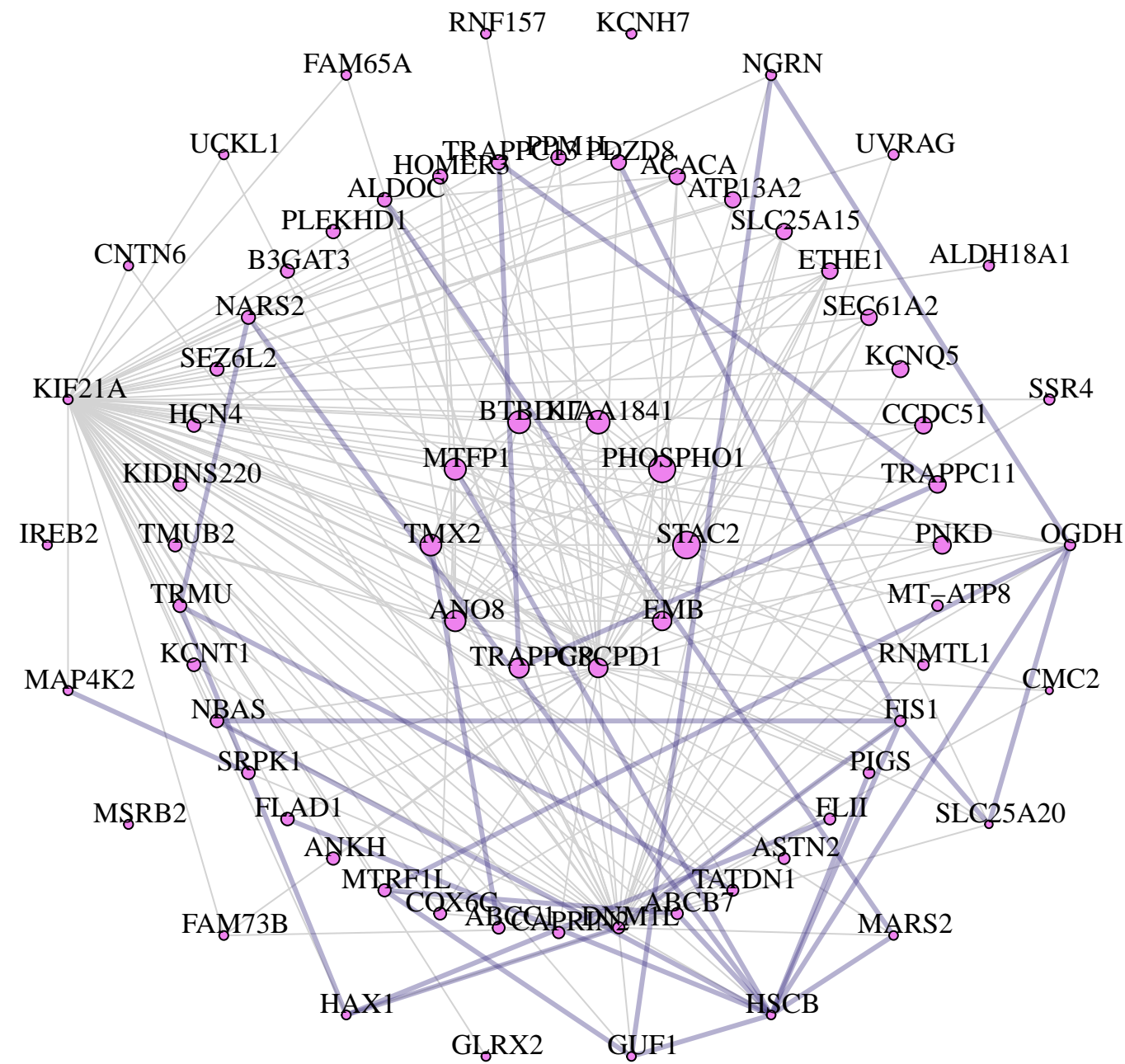


M33 darkolivegreen module hubs connected by top 320 TOM edges: HUB<sup>degree</sup>

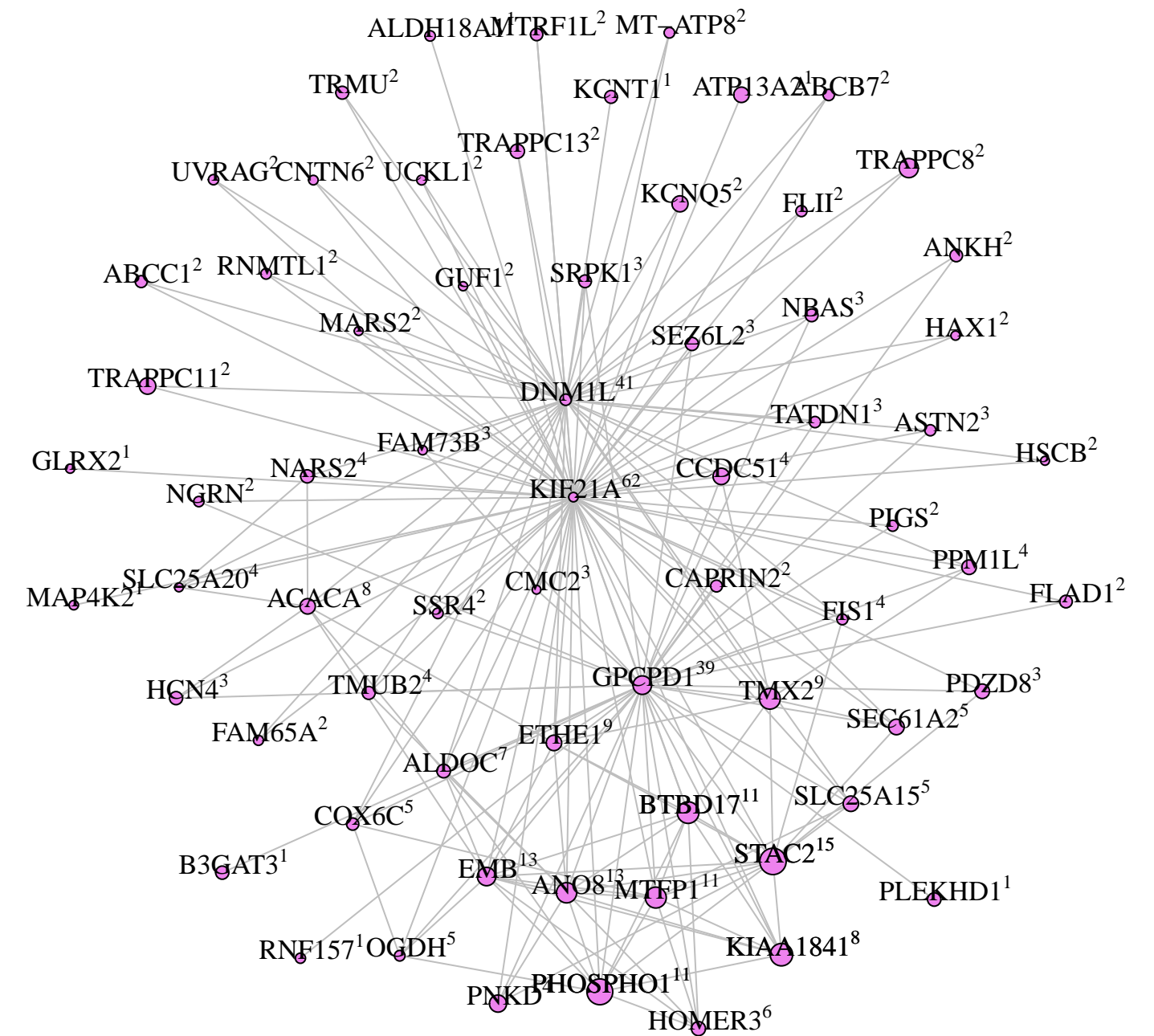




### M32 violet module



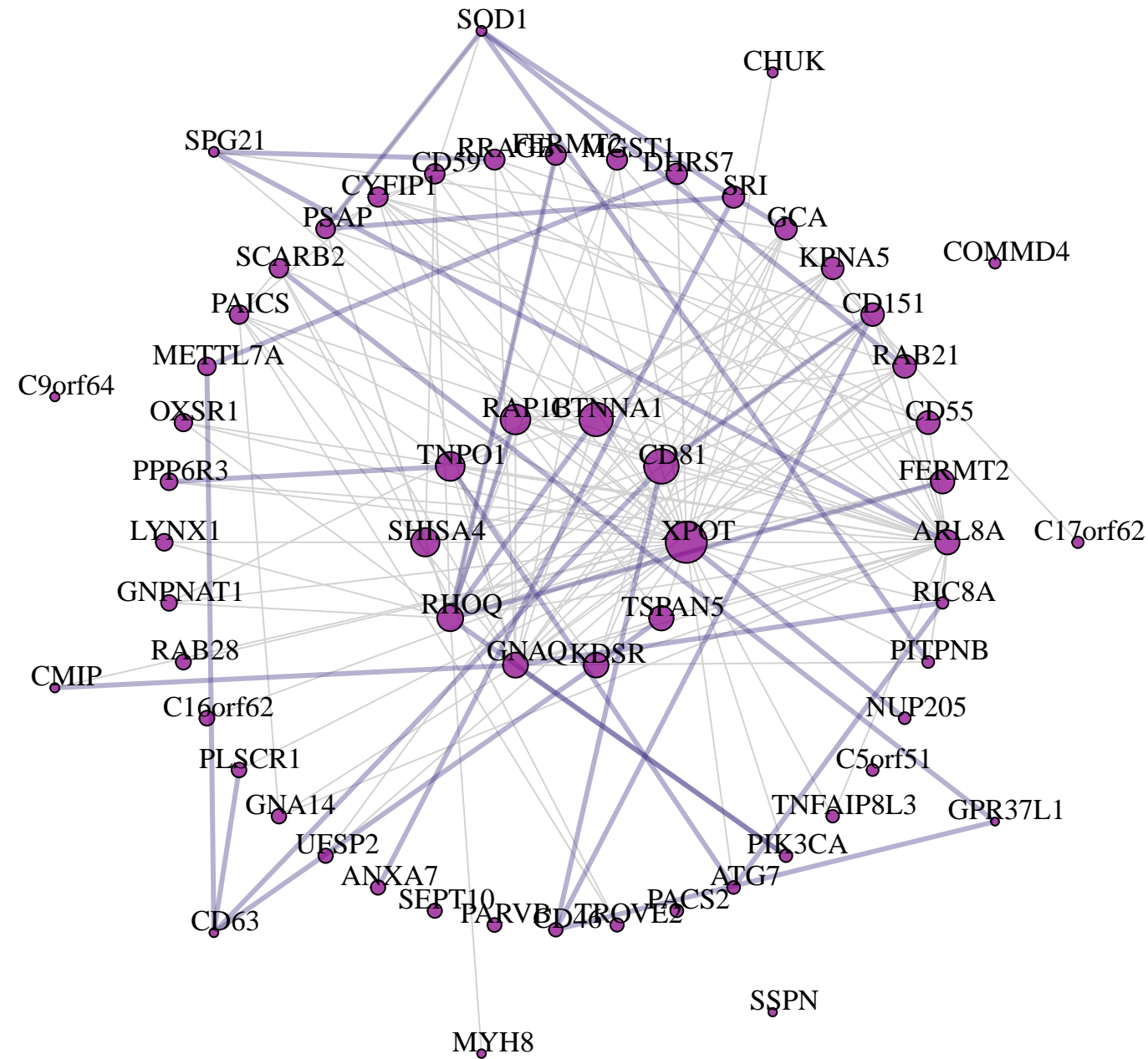
M32 violet module hubs connected by top 396 TOM edges: HUB<sup>degree</sup>



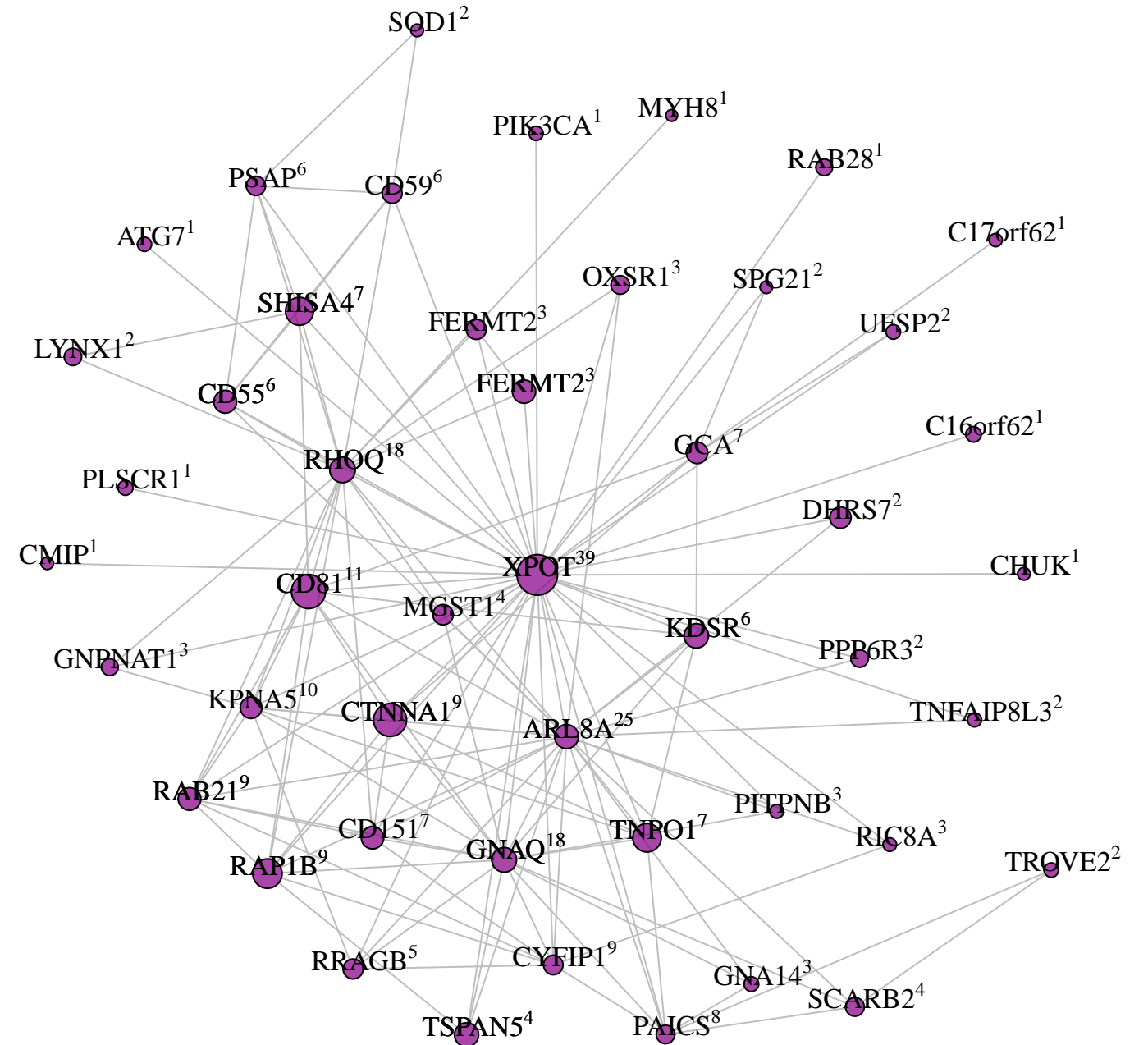
[illegible]



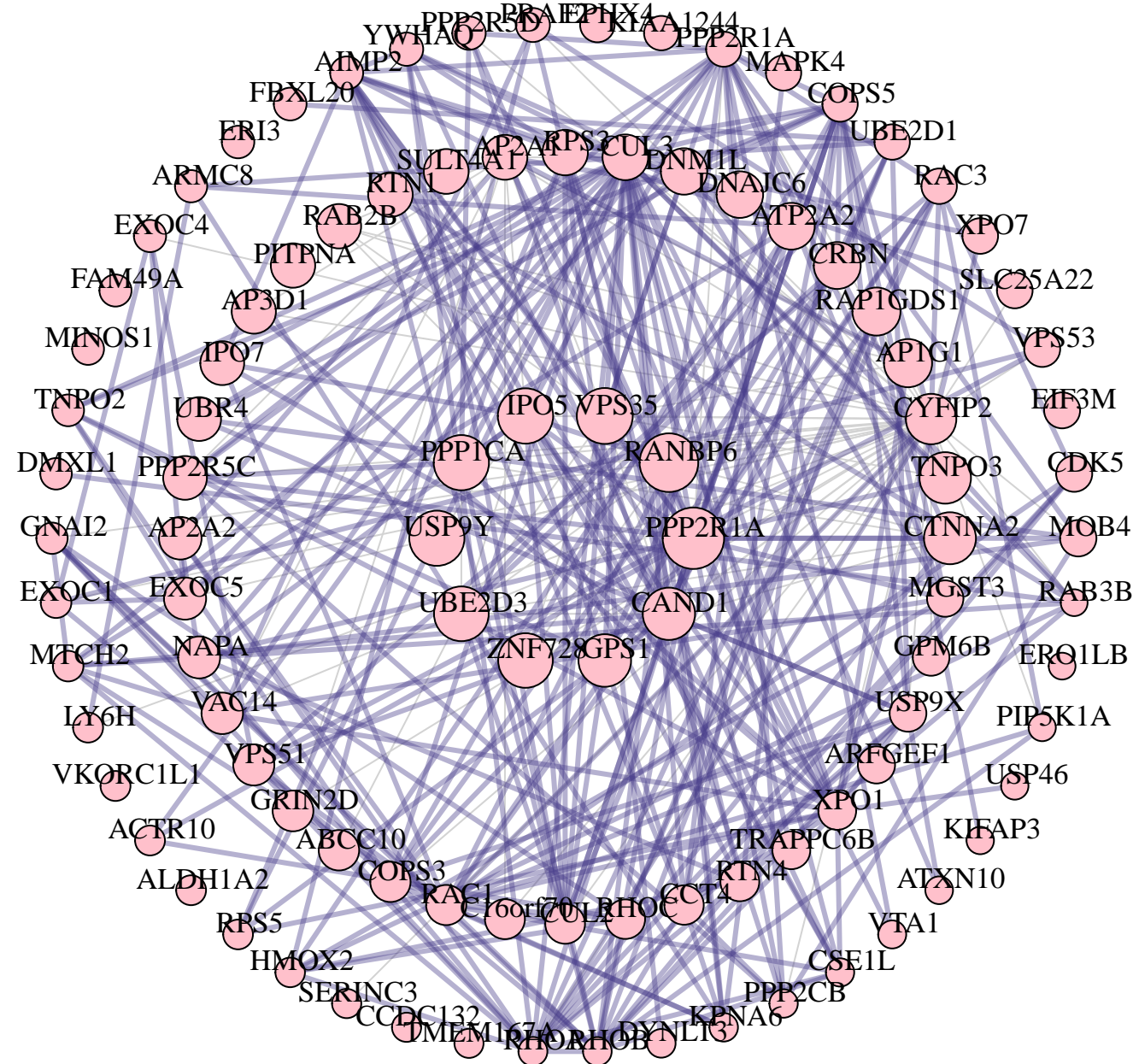
M34 darkmagenta module



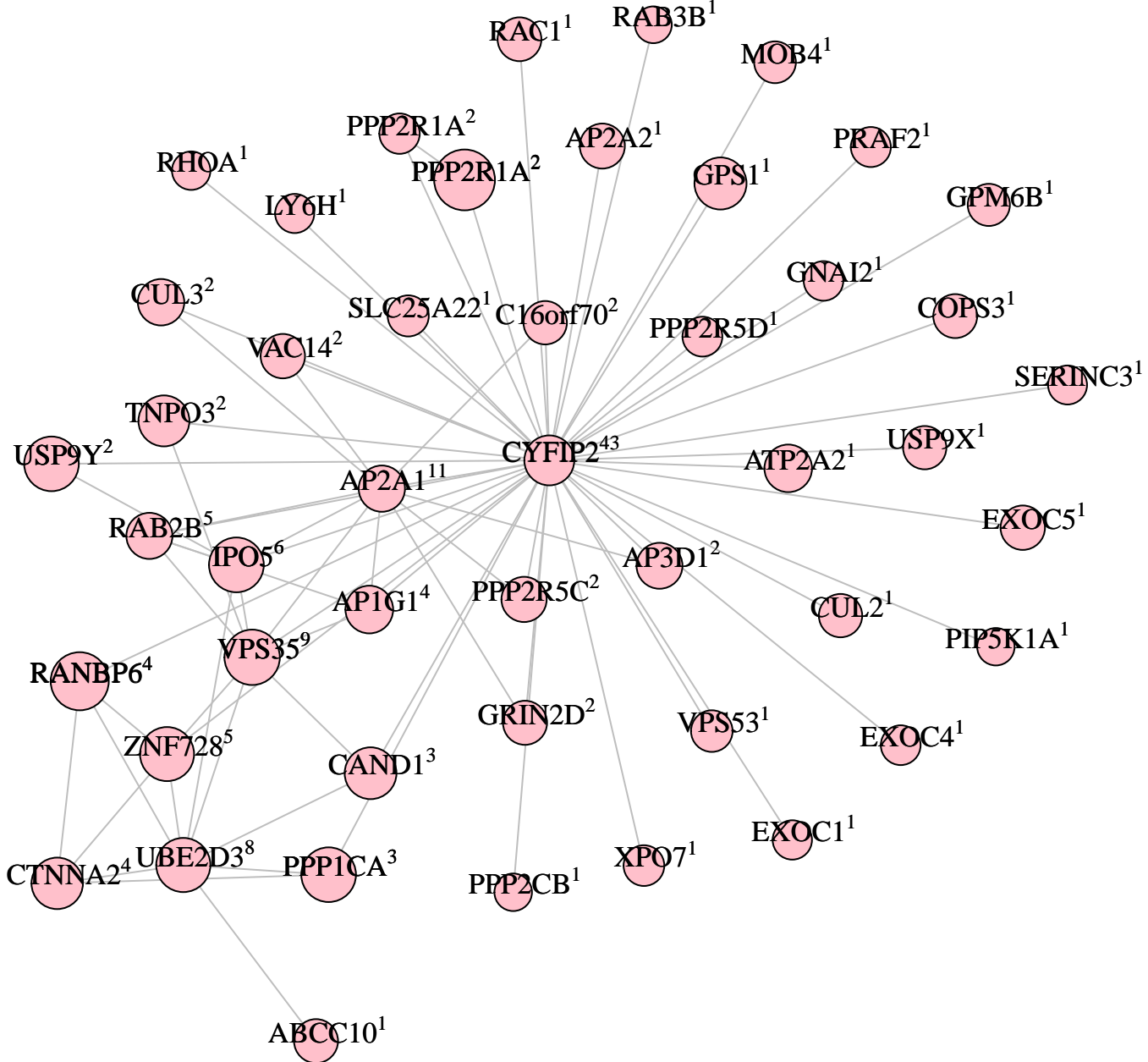
M34 darkmagenta module hubs connected by top 280 TOM edges: HUB<sup>degree</sup>



M8 pink module



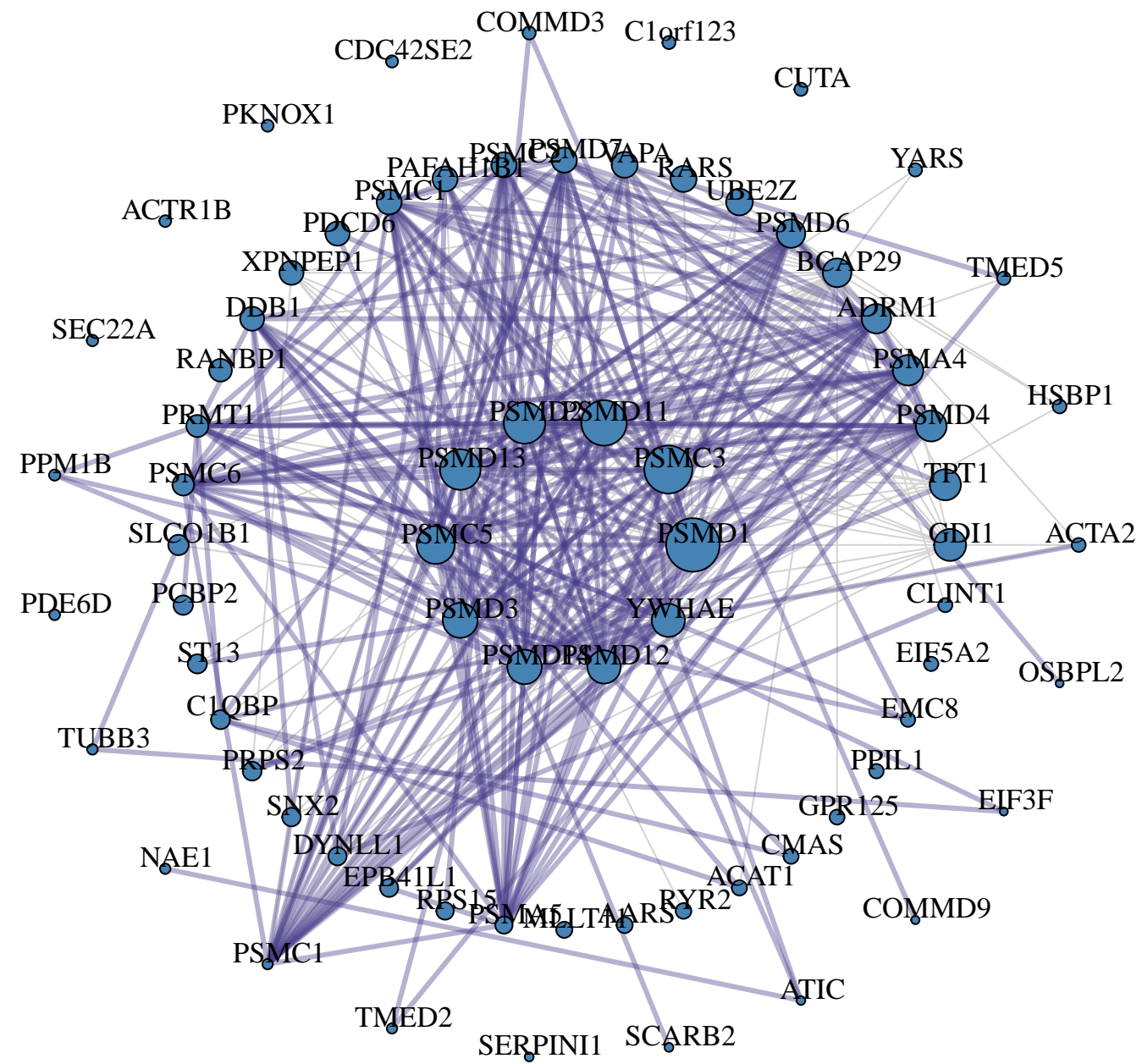
M8 pink module hubs connected by top 150 TOM edges: HUB<sup>degree</sup>



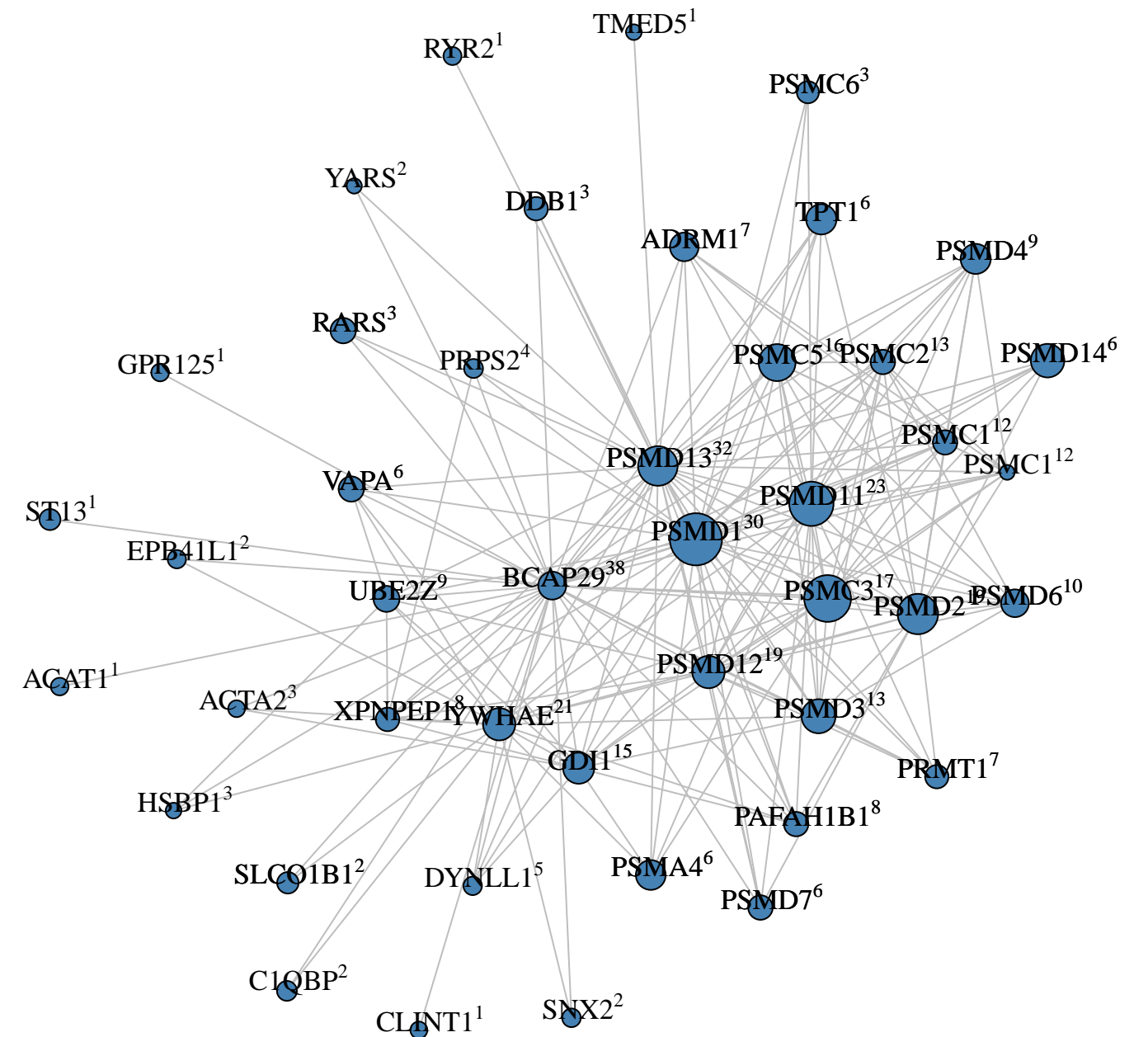




M30 steelblue module

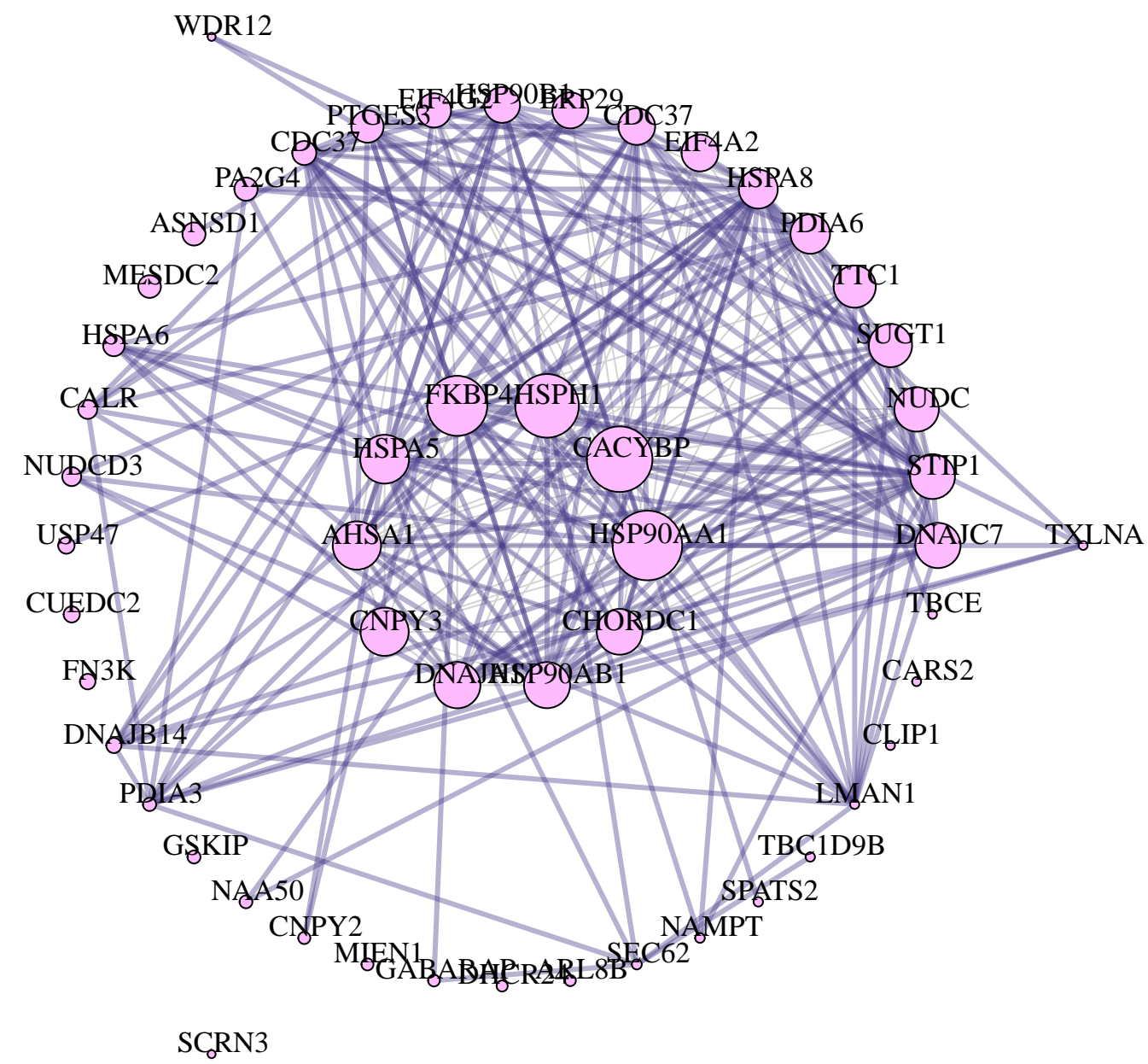


M30 steelblue module hubs connected by top 408 TOM edges: HUB<sup>degree</sup>

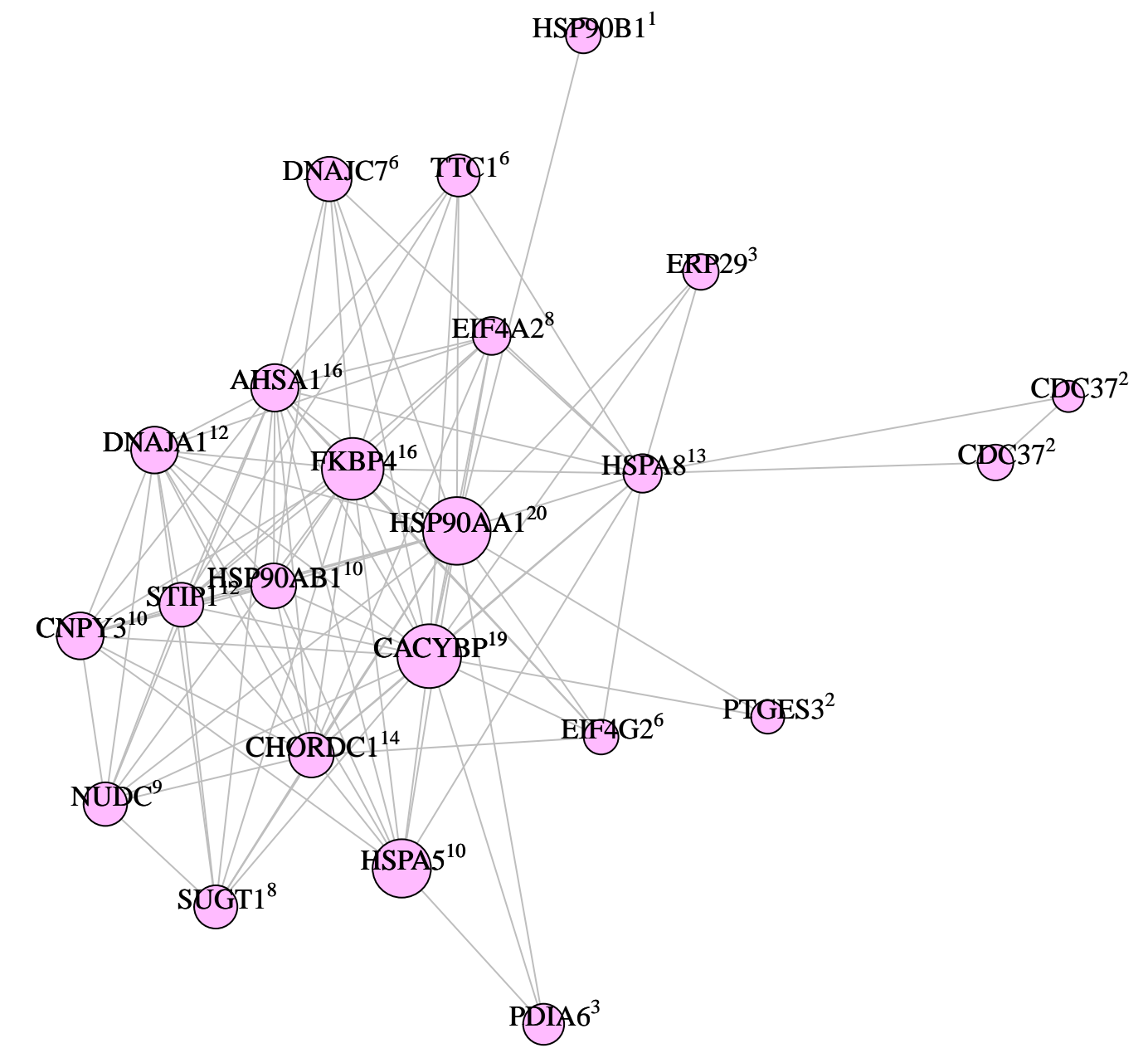




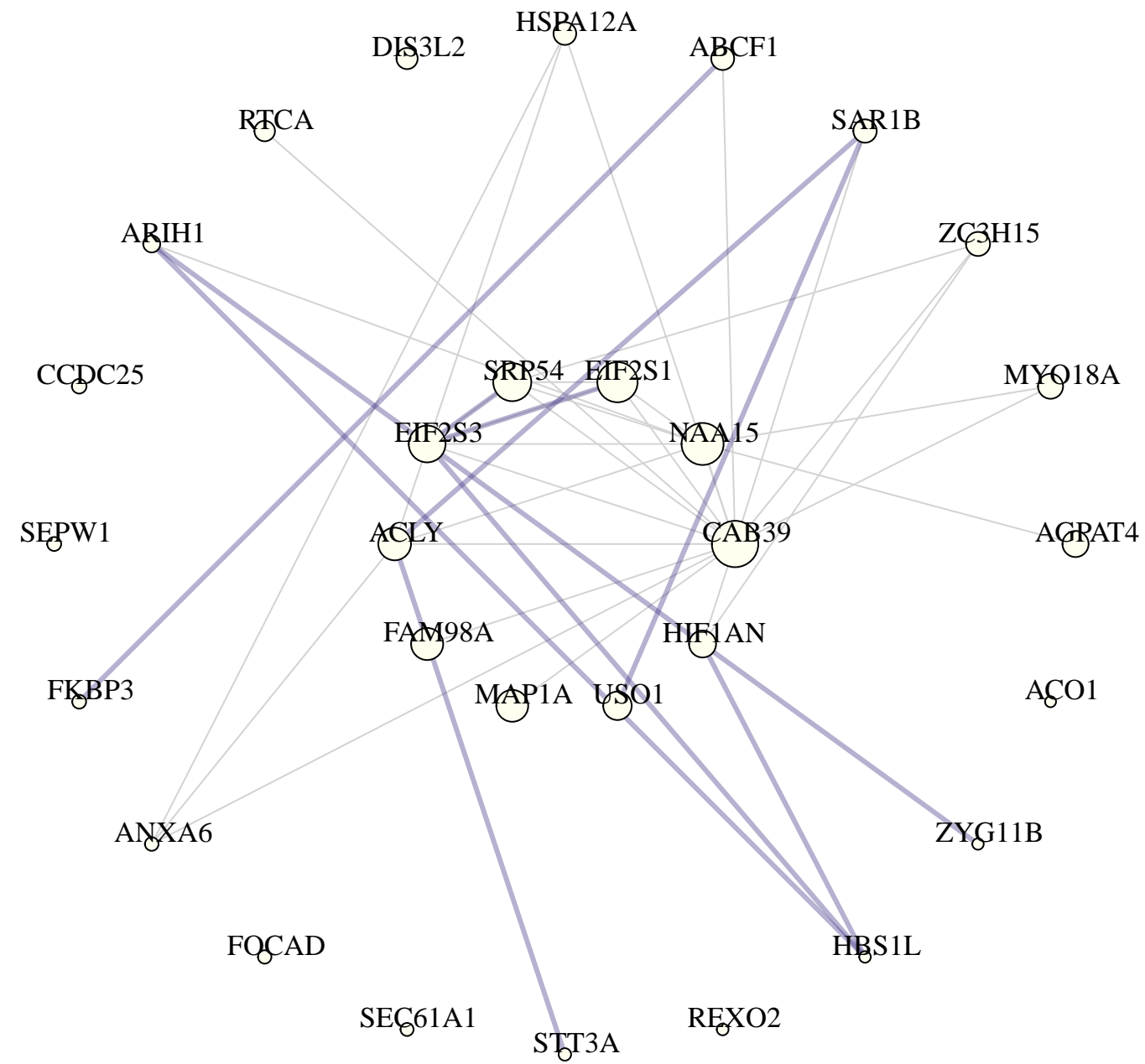
**M38 plum1 module**



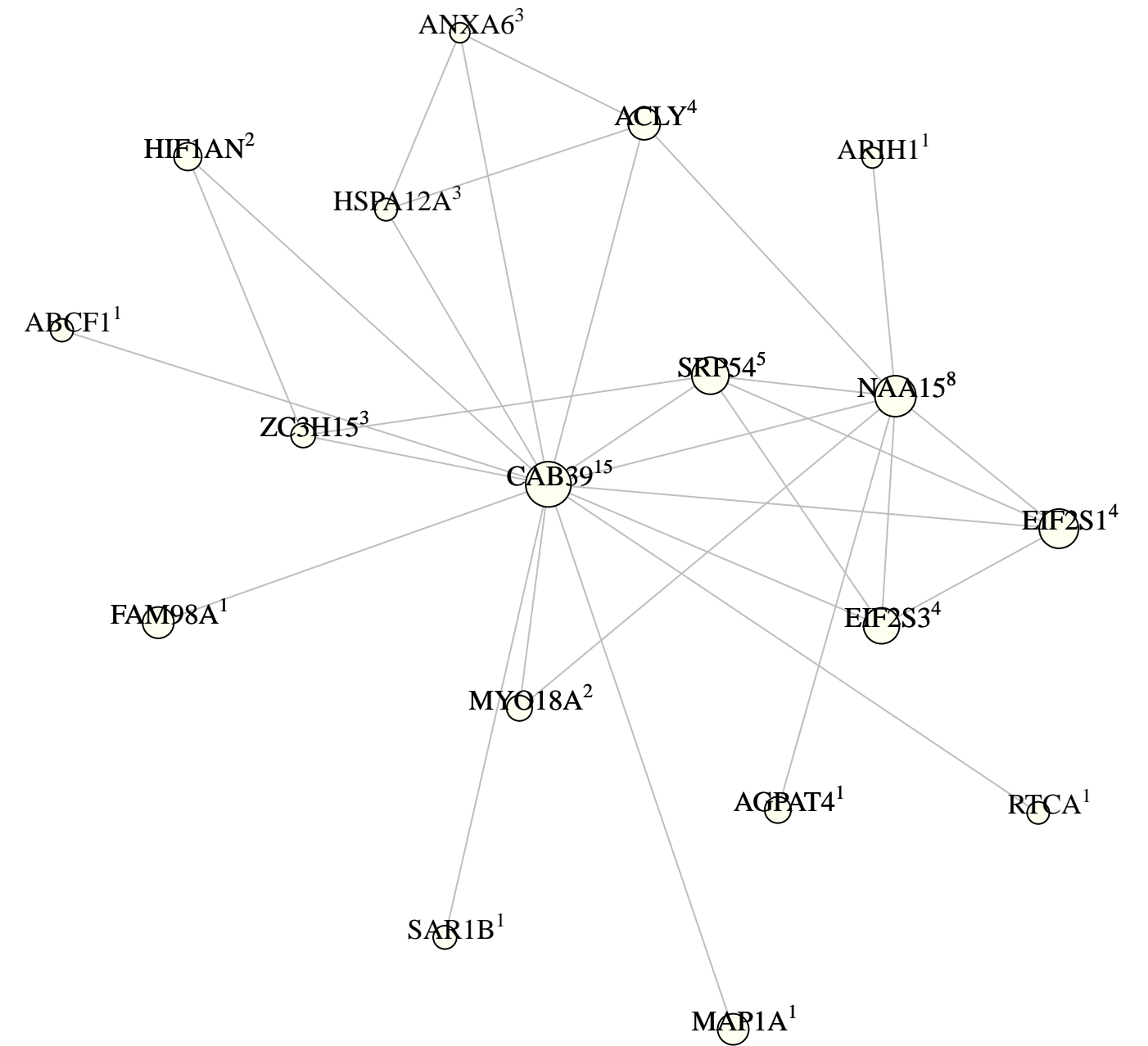
M38 plum1 module hubs connected by top 208 TOM edges: HUB<sup>degree</sup>



**M43 ivory module**

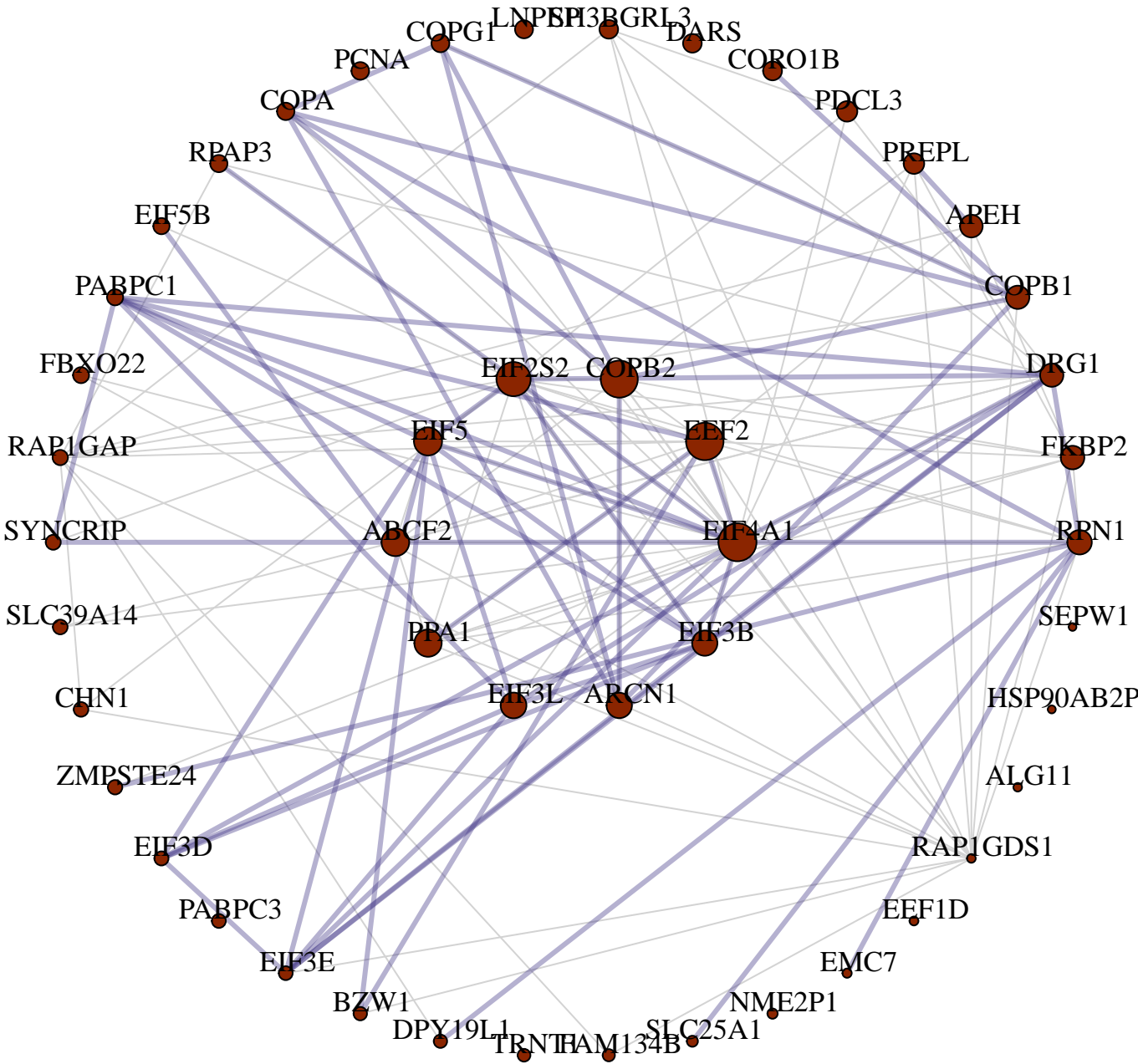


M43 ivory module hubs connected by top 60 TOM edges: HUB<sup>degree</sup>

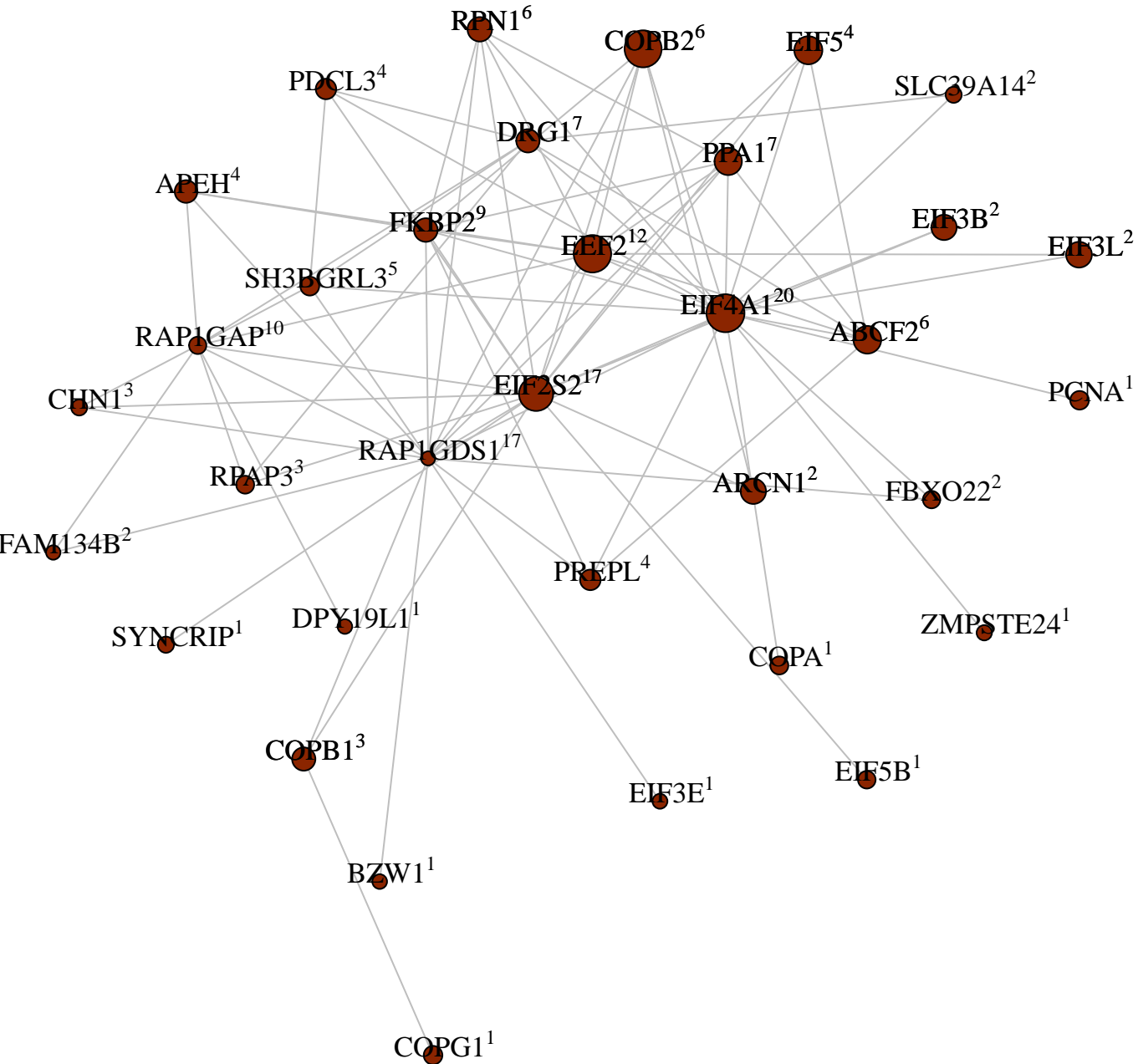




M39 orangered4 module



M39 orangered4 module hubs connected by top 168 TOM edges: HUB<sup>degree</sup>



SYN1<sup>47</sup>

SCYL1<sup>1</sup>

RASA1<sup>1</sup>

PPME1<sup>1</sup>

UBE2N<sup>1</sup>

SAR1A<sup>1</sup>

SACM1L<sup>1</sup>

DYNC1H1<sup>1</sup>

TRAPPC3<sup>1</sup>

PPP2R2D<sup>1</sup>

STAM<sup>1</sup>

RAN<sup>1</sup>

HSPA4<sup>2</sup>

LDHA<sup>1</sup>

CCT2<sup>4</sup>

CCT5<sup>9</sup>

CCT8<sup>10</sup>

USP14<sup>2</sup>

PFKP<sup>1</sup>

AP1S1<sup>2</sup>

RABGEF1<sup>1</sup>

CAPZB<sup>2</sup>

OTUB1<sup>1</sup>

RABEP1<sup>1</sup>

RAP1GAP<sup>1</sup>

ATP2C1<sup>1</sup>

RAB7A<sup>1</sup>

DSTN<sup>1</sup>

GMFB<sup>1</sup>

AP1M2<sup>1</sup>

PPP5C<sup>1</sup>

PPP1R7<sup>1</sup>

HAGH<sup>1</sup>

NAPG<sup>1</sup>

DUSP3<sup>1</sup>

UBE2M<sup>1</sup>

ATG5<sup>1</sup>

TMEM189

UBE2V1<sup>1</sup>

COPS4<sup>1</sup>

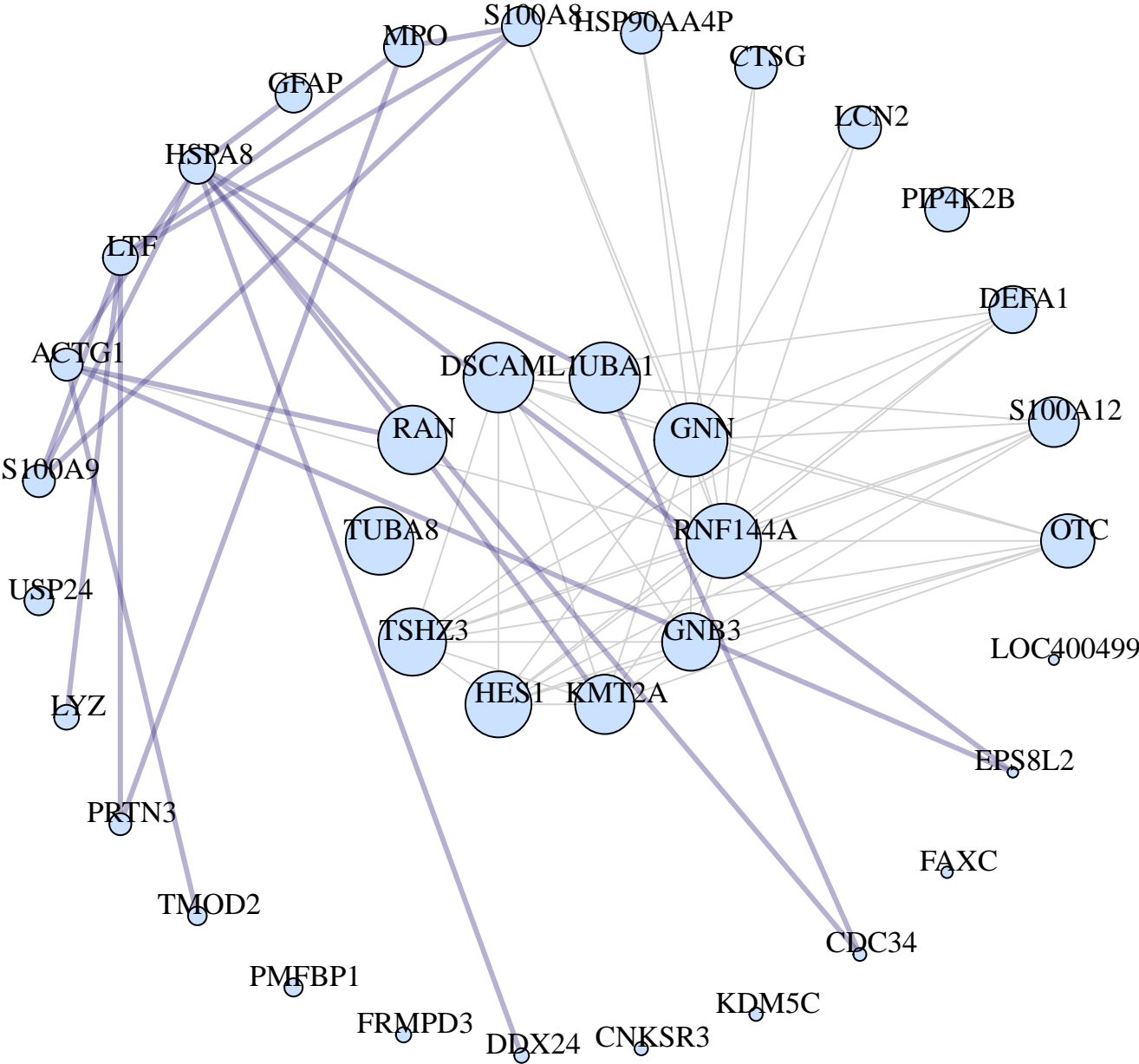
DNAJA2<sup>1</sup>

DYNC1LI1<sup>1</sup>

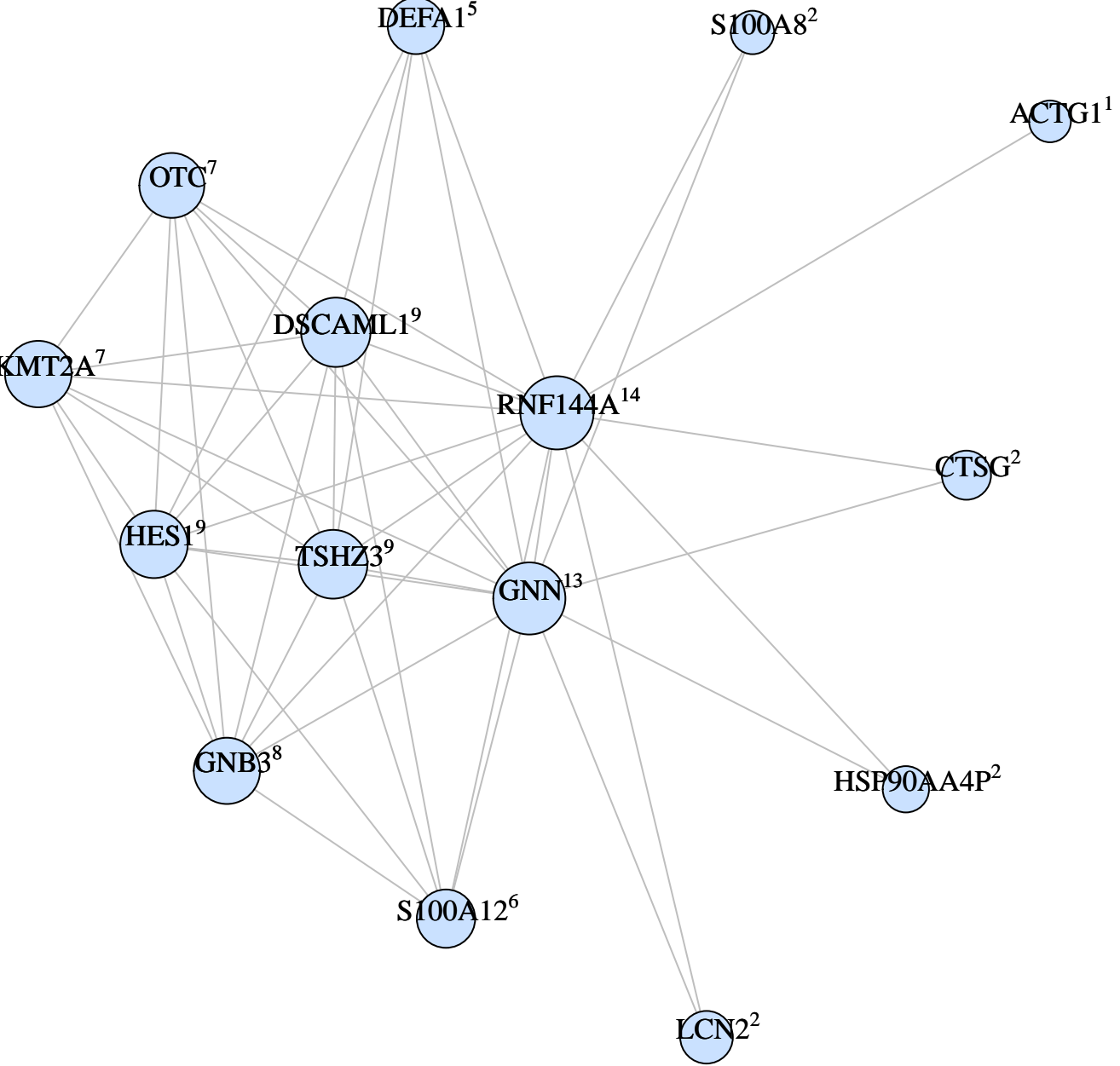


Protein-Protein Interactions (PPIs) network diagram showing interactions between various proteins. The nodes are labeled with protein names and their associated counts (e.g., DYNC1H1<sup>15</sup>, CAPN1<sup>6</sup>, CAPNS1<sup>5</sup>, NRIP2<sup>4</sup>, etc.). The network is highly interconnected, with many nodes having multiple connections.

M41 lightsteelblue1 module

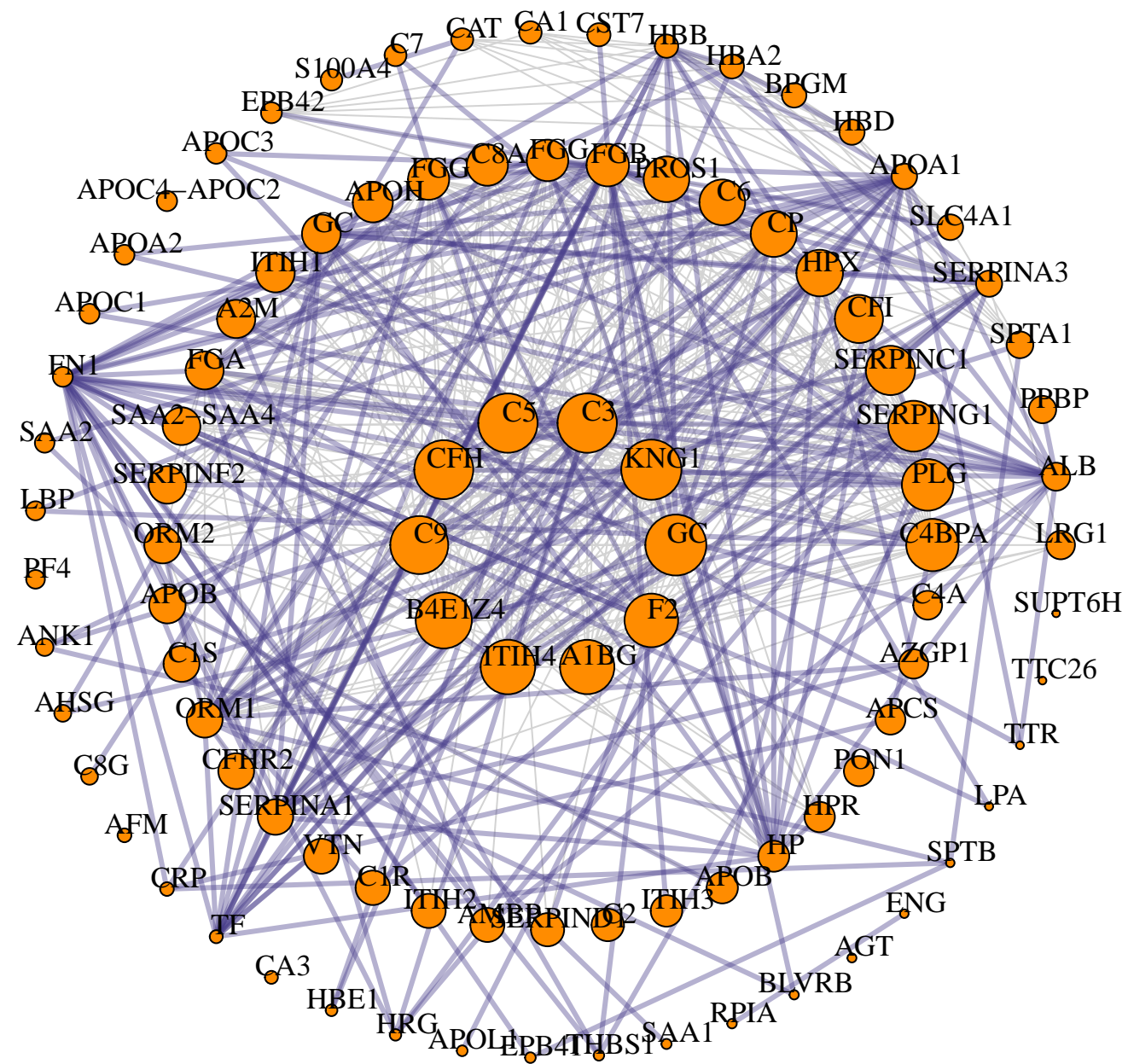


M41 lightsteelblue1 module hubs connected by top 96 TOM edges: HUB<sup>degree</sup>

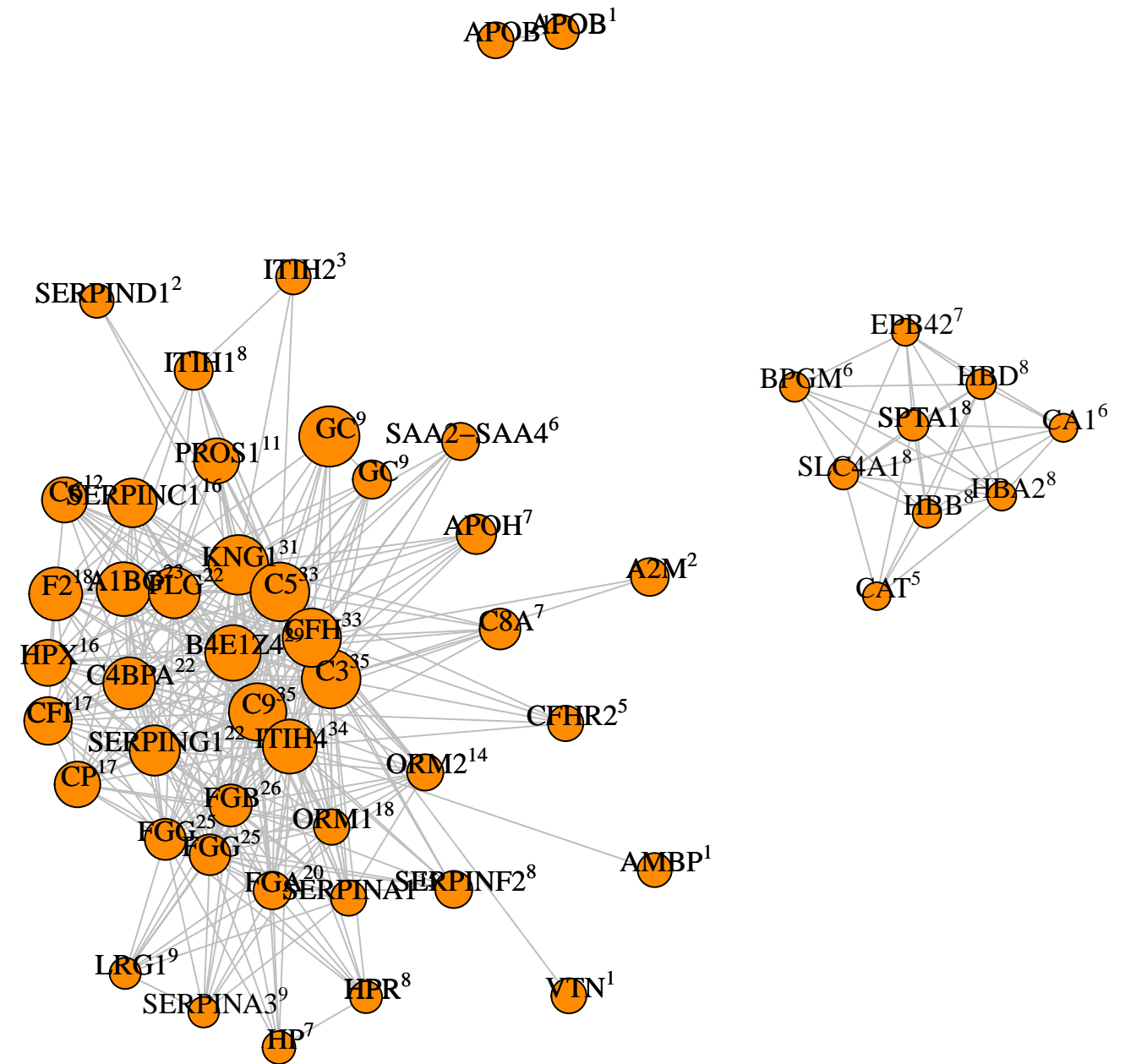




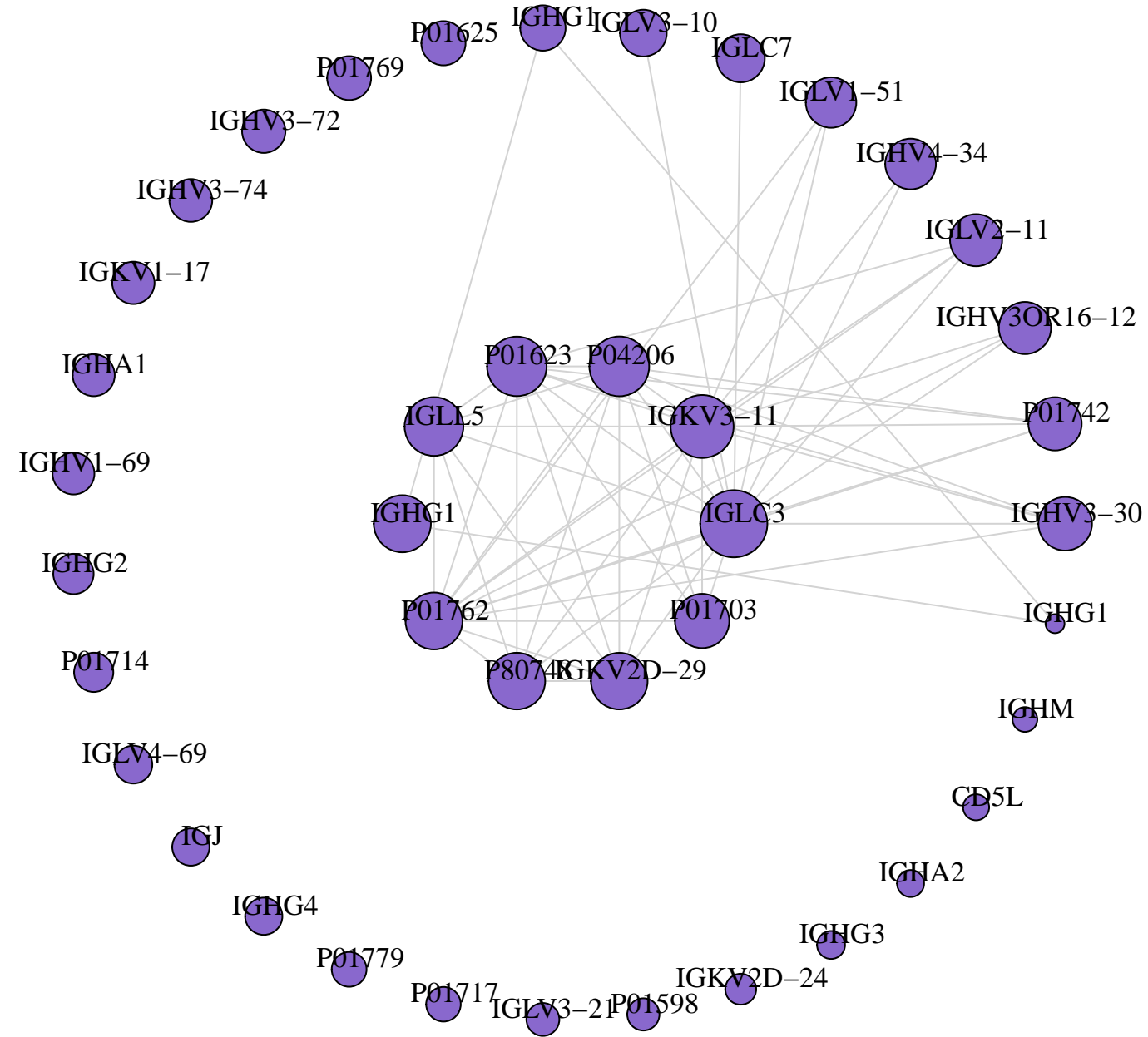
## M26 darkorange module



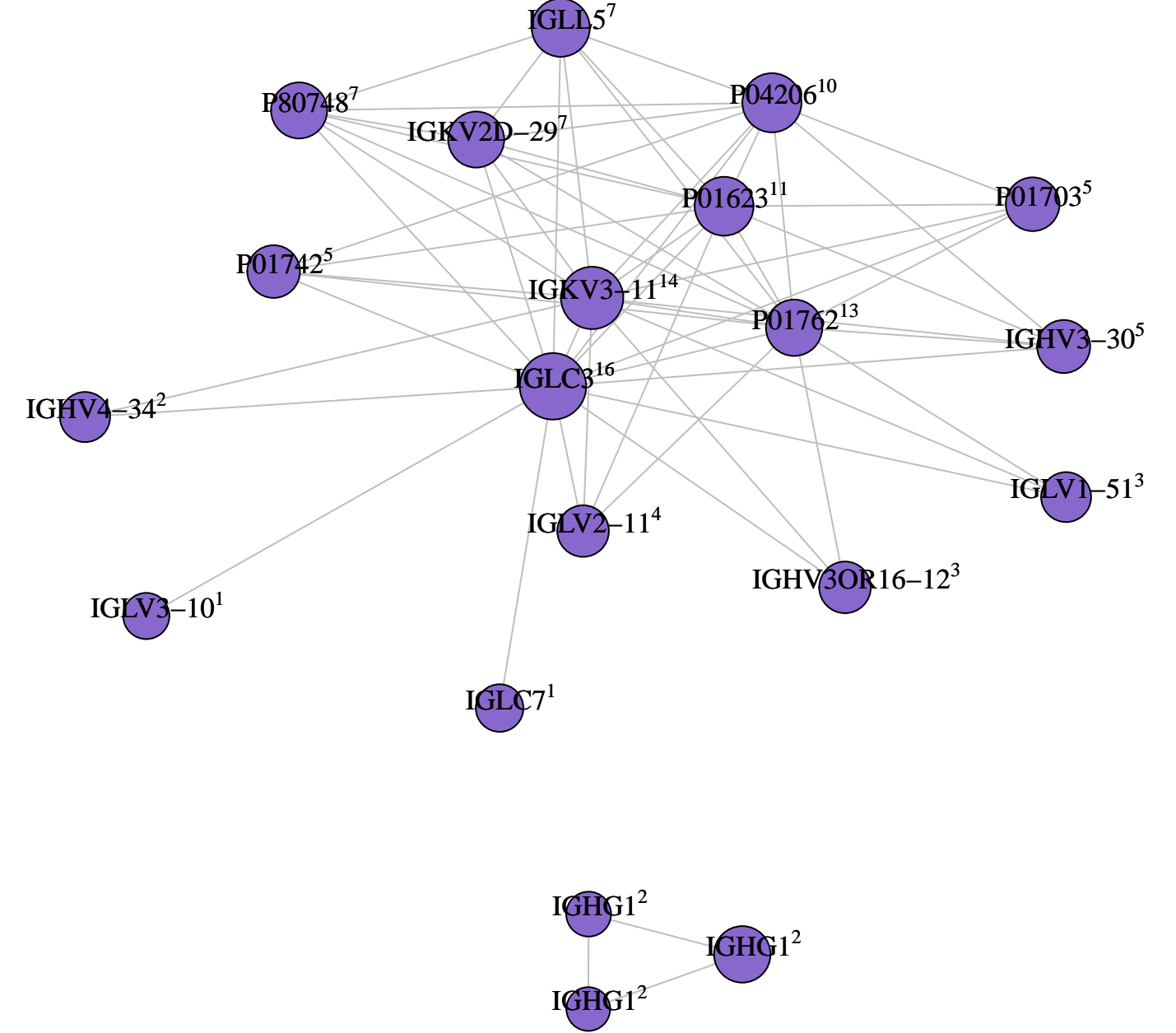
M26 darkorange module hubs connected by top 736 TOM edges: HUB<sup>degree</sup>



## M40 mediumpurple3 module

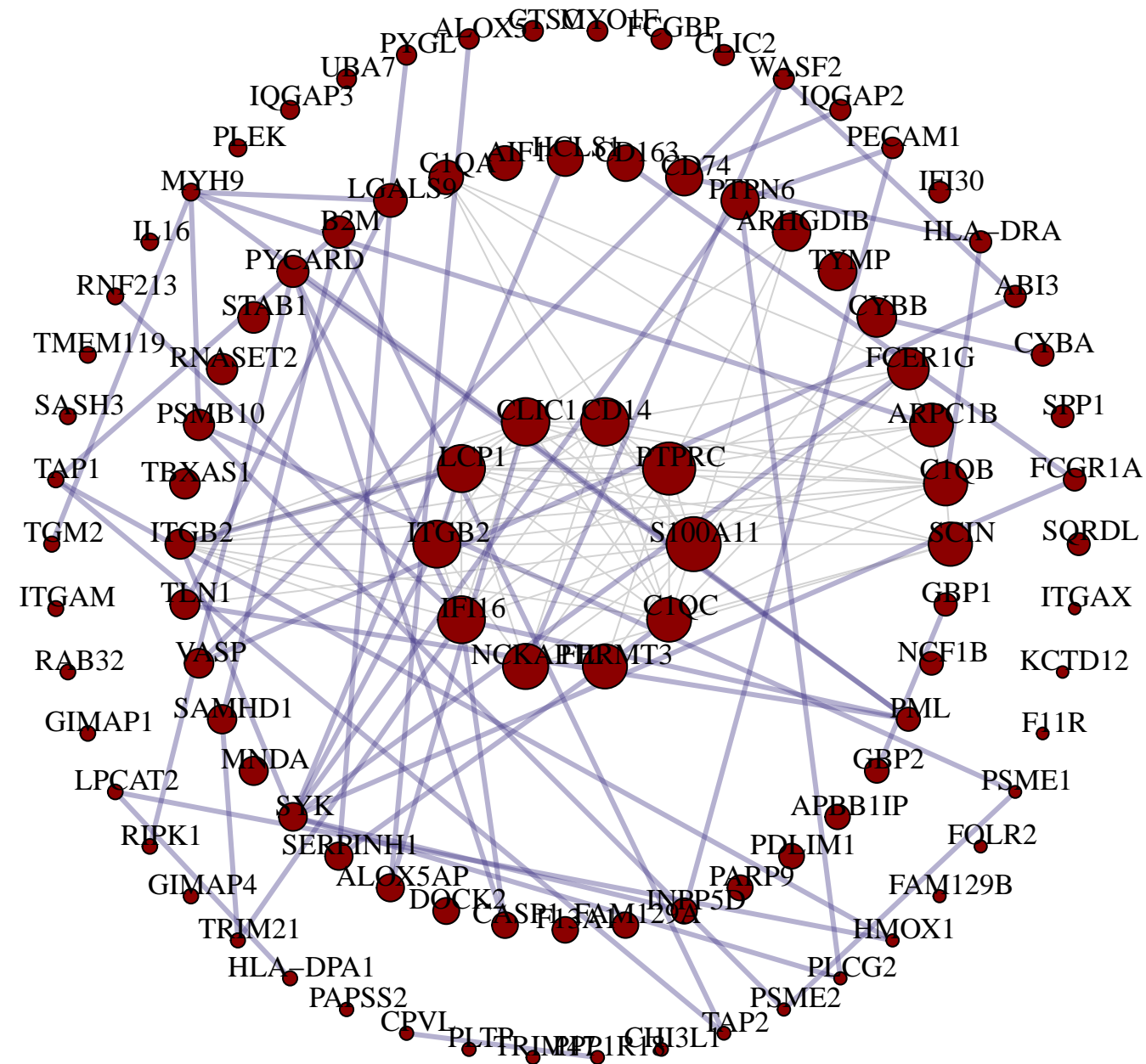


M40 mediumpurple3 module hubs connected by top 120 TOM edges: HUB<sup>degree</sup>

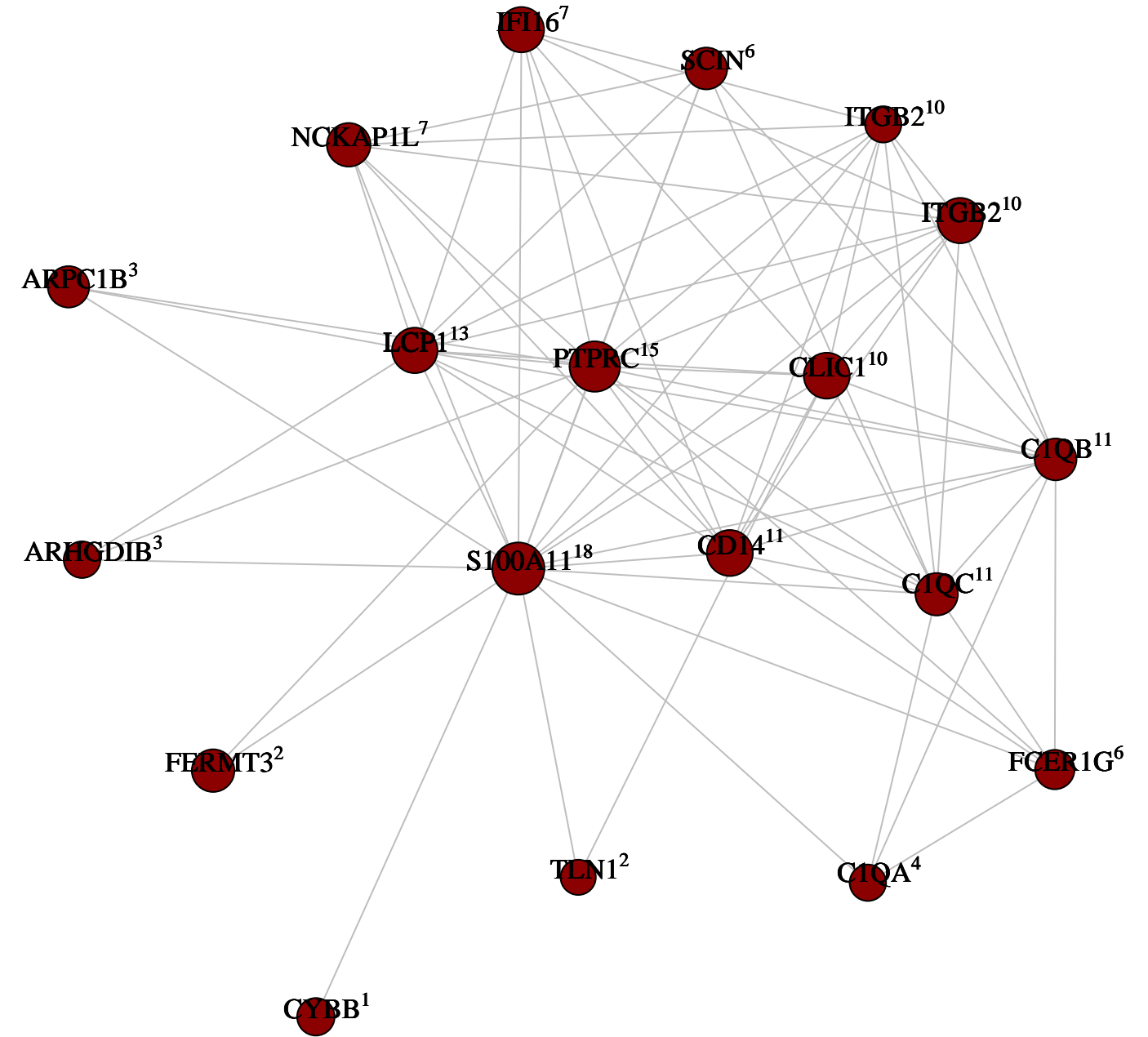




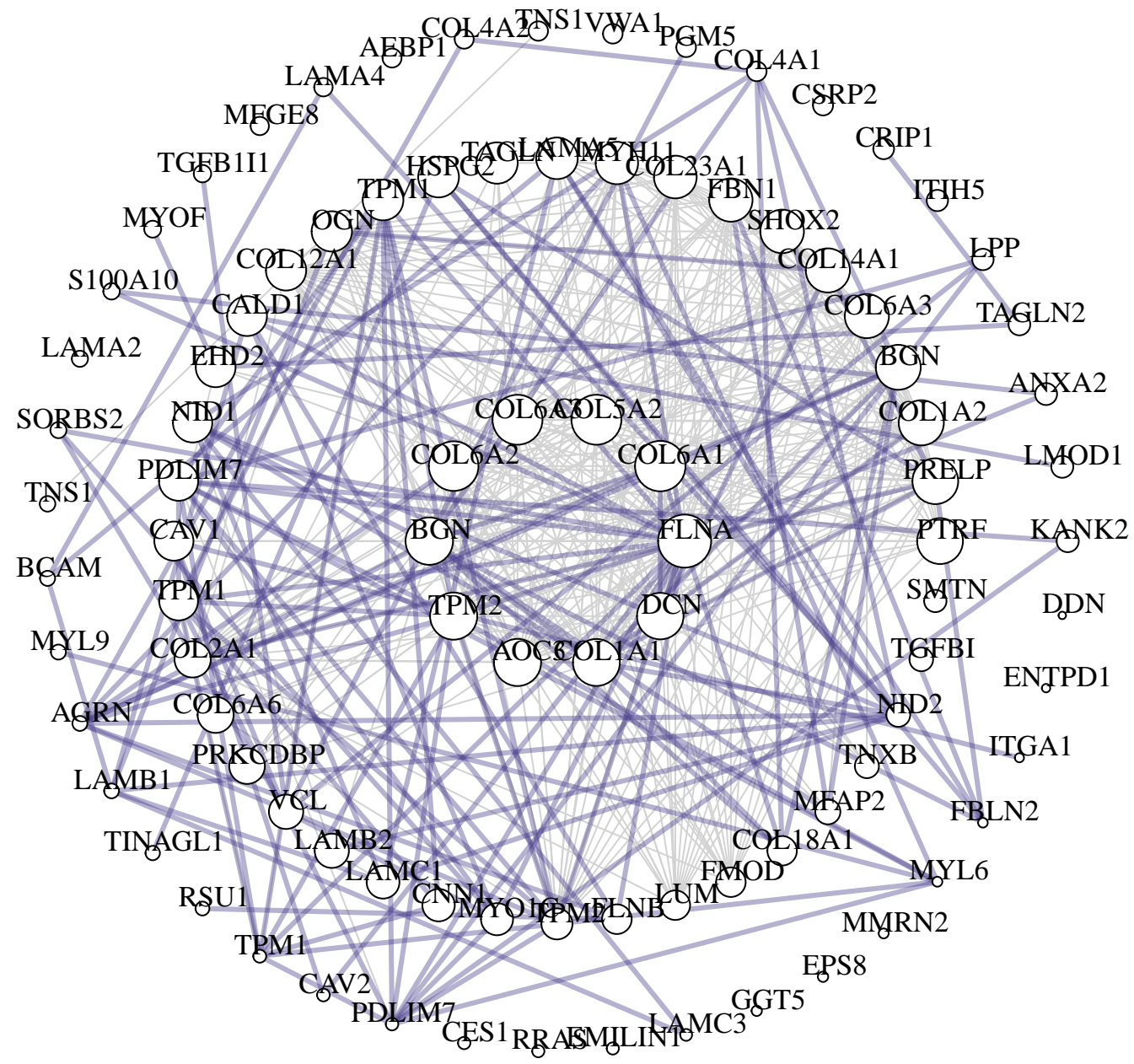
M21 darkred module



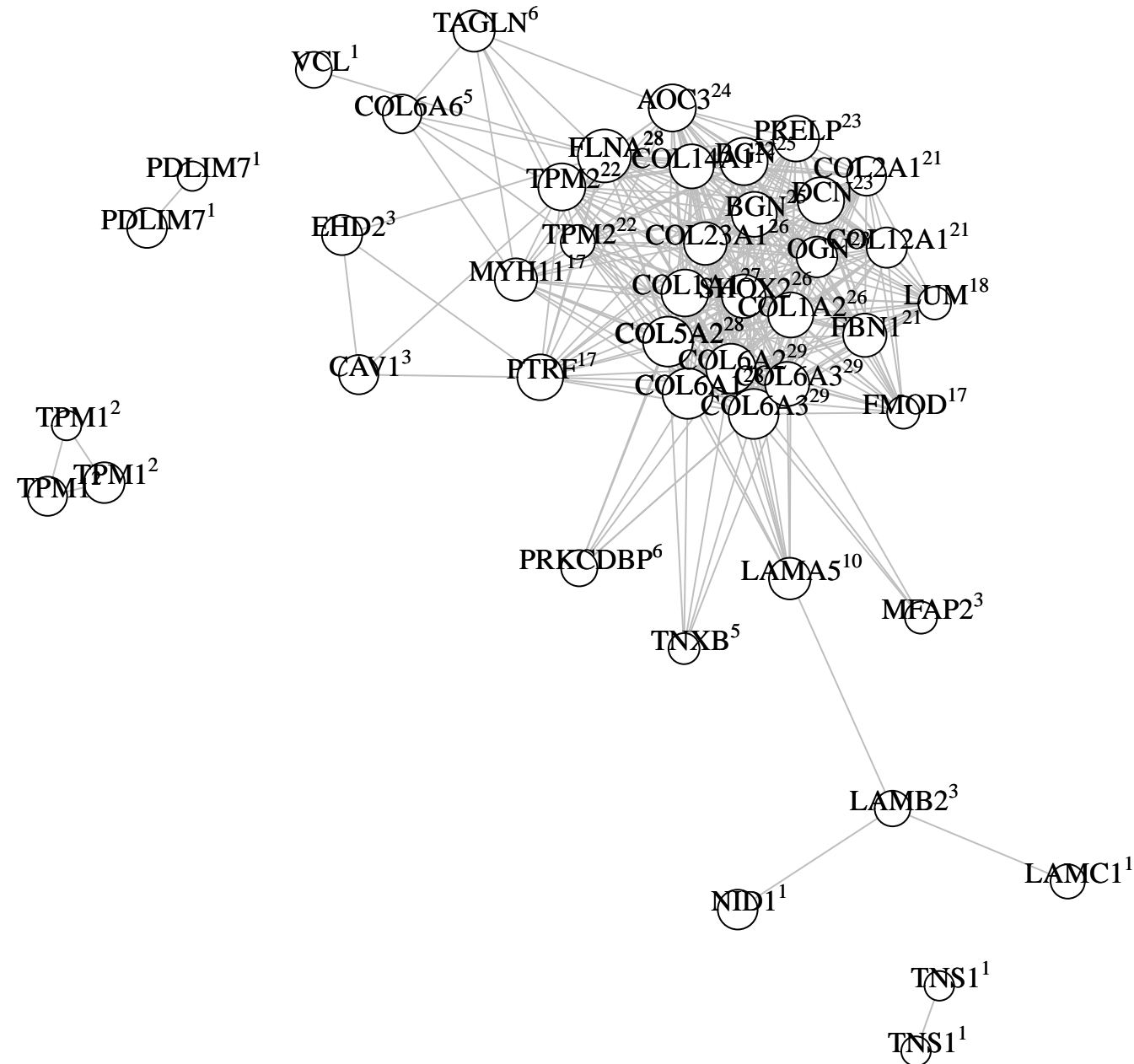
M21 darkred module hubs connected by top 150 TOM edges: HUB<sup>degree</sup>



M27 white module



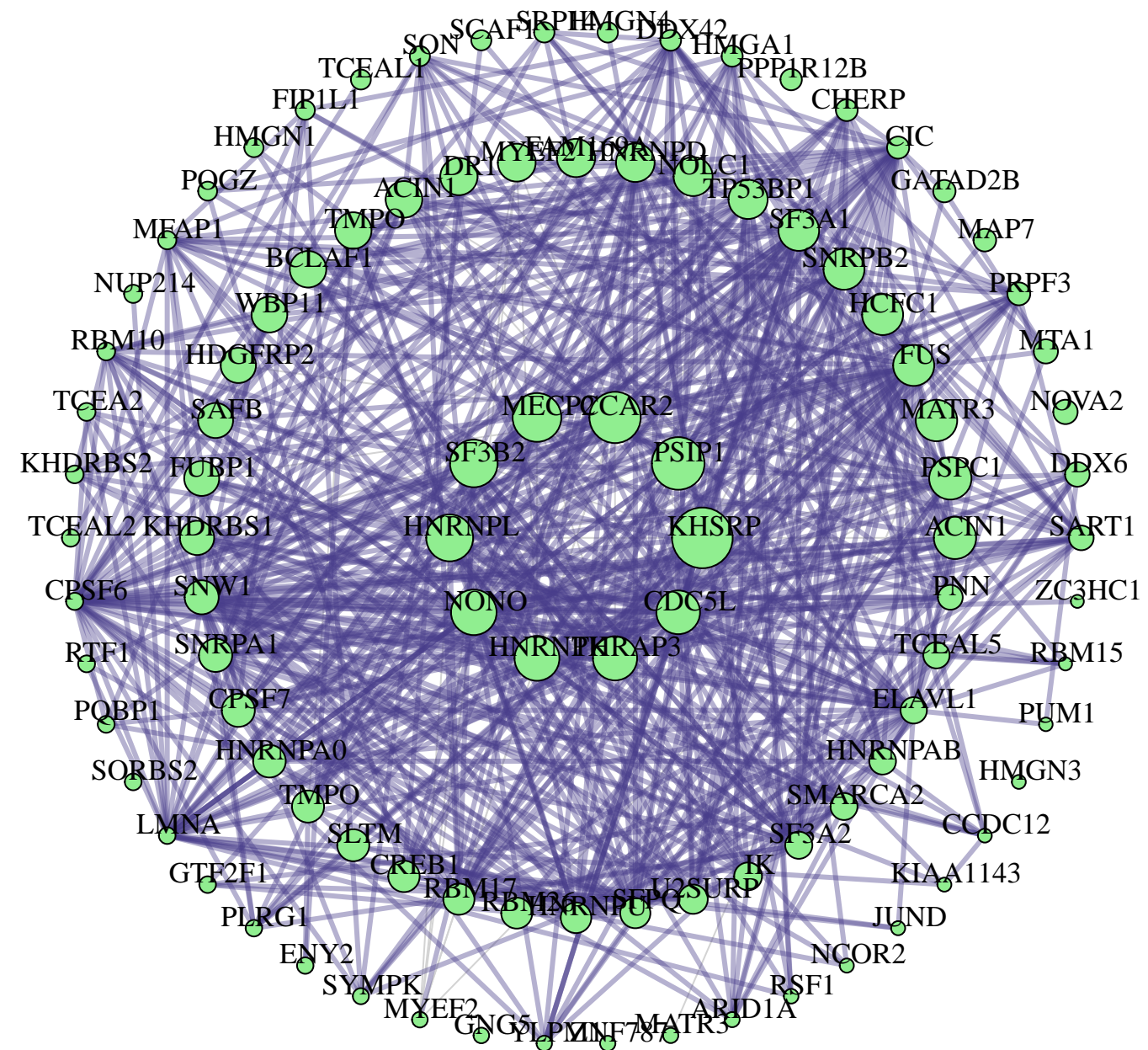
M27 white module hubs connected by top 674 TOM edges: HUB<sup>degree</sup>



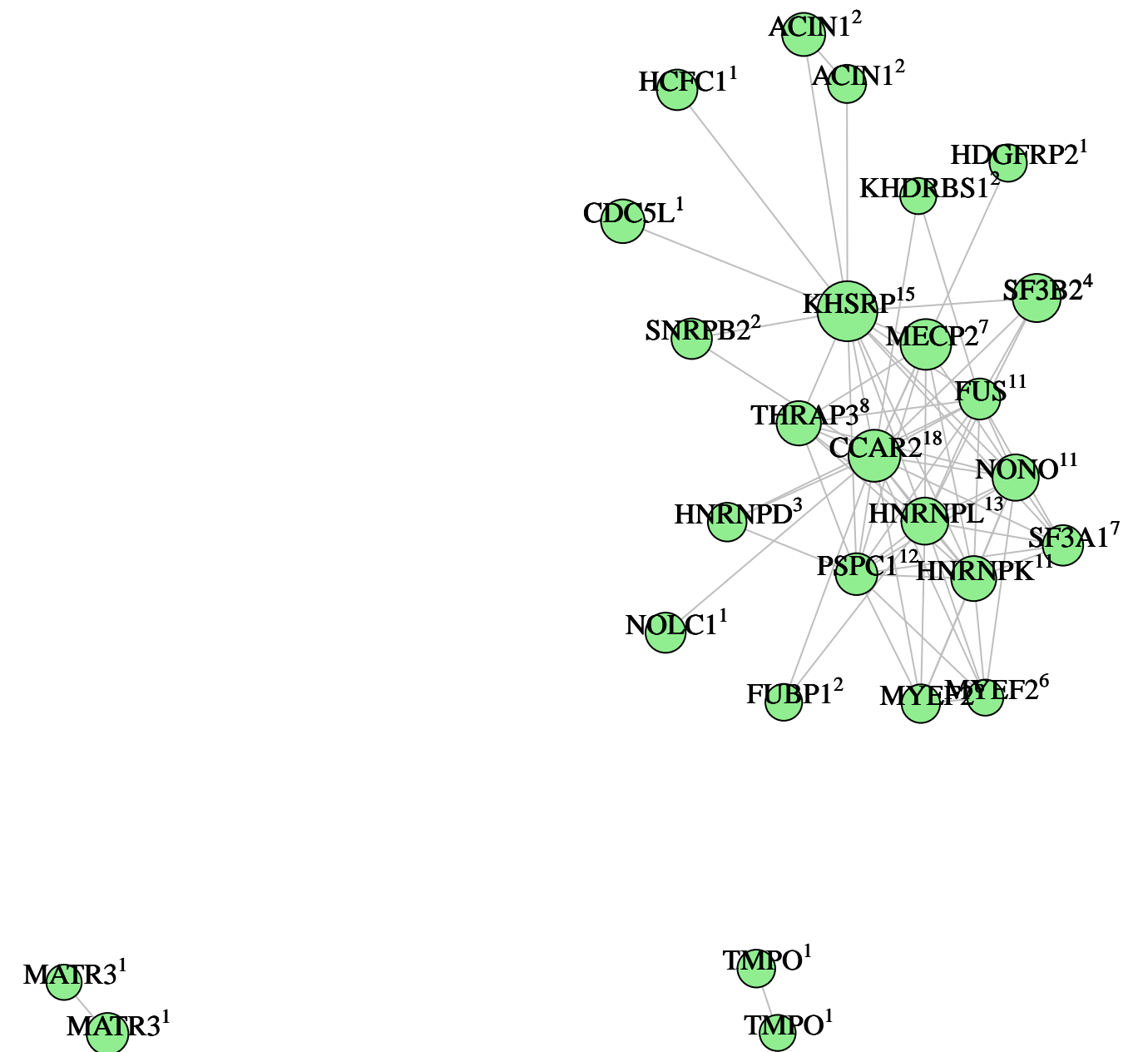


Network diagram showing interactions between 25 genes. Nodes are labeled with gene names and counts in parentheses. The network is highly interconnected, with a central cluster of genes including UBAP2L, H0YHG0, MTDH, ZRANB2, RBMX, SNRPC, SERBP1, HABP4, ATXN2L, and ATXN2. Other genes like NUP153, PPP1R2, LSM14A, and ALYREF are also part of the central cluster. Genes like G3BP2 and G3BP2 are isolated at the bottom left.

## M18 lightgreen module



M18 lightgreen module hubs connected by top 150 TOM edges: HUB<sup>degree</sup>





[illegible]

Network diagram showing interactions between 20 genes. The nodes are labeled with gene names and superscripted numbers. The edges represent interactions between the genes.

Genes and their counts (from top to bottom, left to right):

- DHX9<sup>1</sup>
- RBM8A<sup>2</sup>
- DDX46<sup>2</sup>
- HNRNPK<sup>2</sup>
- HNRNPA2B1<sup>8</sup>
- DDX17<sup>12</sup>
- RBM14<sup>3</sup>
- CBX5<sup>5</sup>
- EDF1<sup>4</sup>
- SRRT<sup>9</sup>
- RBM4B<sup>7</sup>
- AKAP8L<sup>6</sup>
- C7orf55-LUC7L2<sup>5</sup>
- LUC7L3<sup>6</sup>
- C14orf166<sup>2</sup>
- RBBP7<sup>13</sup>
- LUC7L<sup>20</sup>
- HNRNPH3<sup>3</sup>
- CFDP1<sup>2</sup>
- RBM39<sup>1</sup>
- HTATSF1<sup>1</sup>
- ZMAT2<sup>1</sup>
- SATB2<sup>1</sup>
- PRPF38B<sup>1</sup>
- SMC1A<sup>3</sup>
- RAD21<sup>1</sup>
- PDS5B<sup>6</sup>
- DEK<sup>7</sup>
- DDX1<sup>4</sup>
- PRPF4B<sup>1</sup>
- DHX15<sup>8</sup>

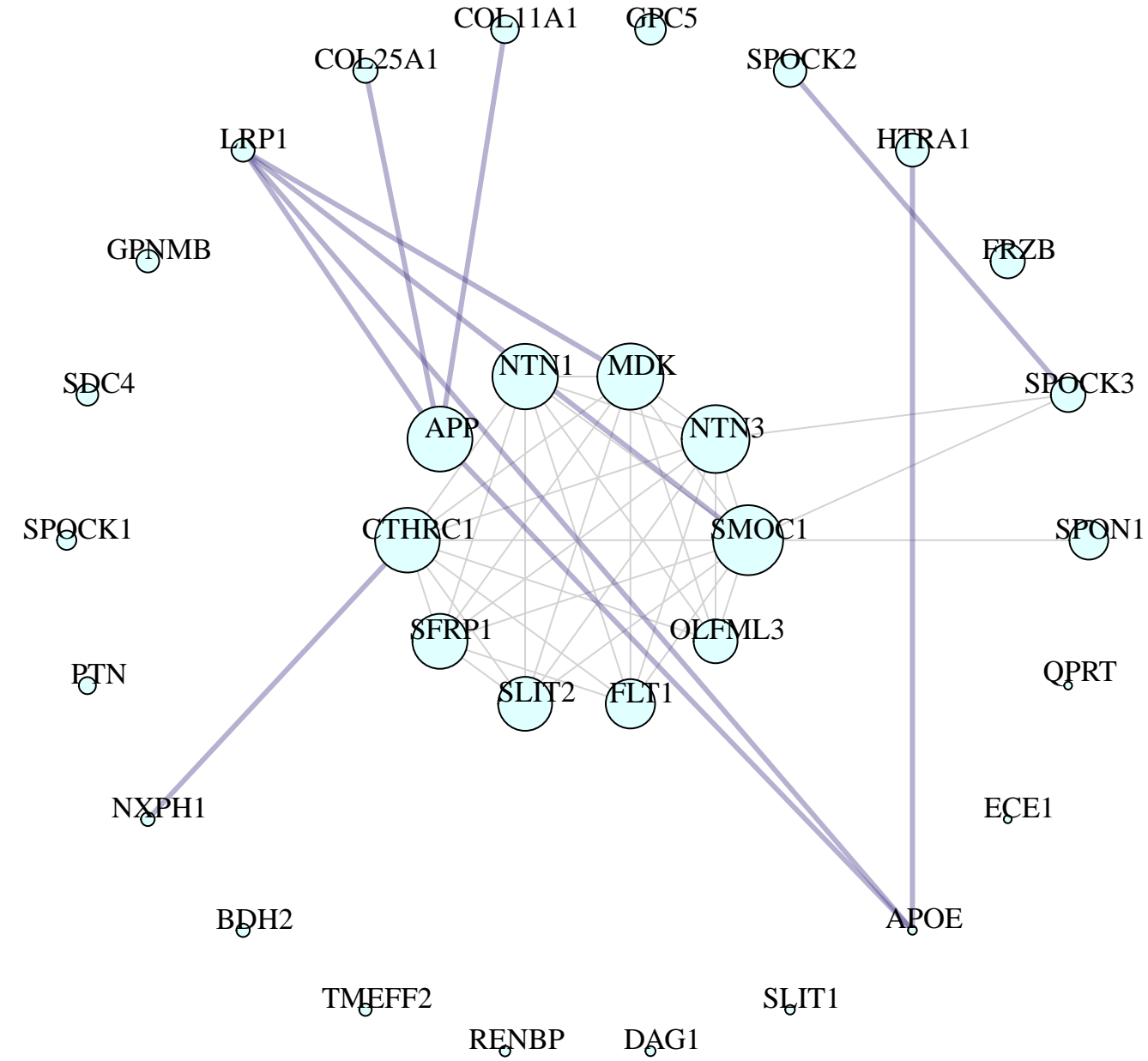


[illegible]

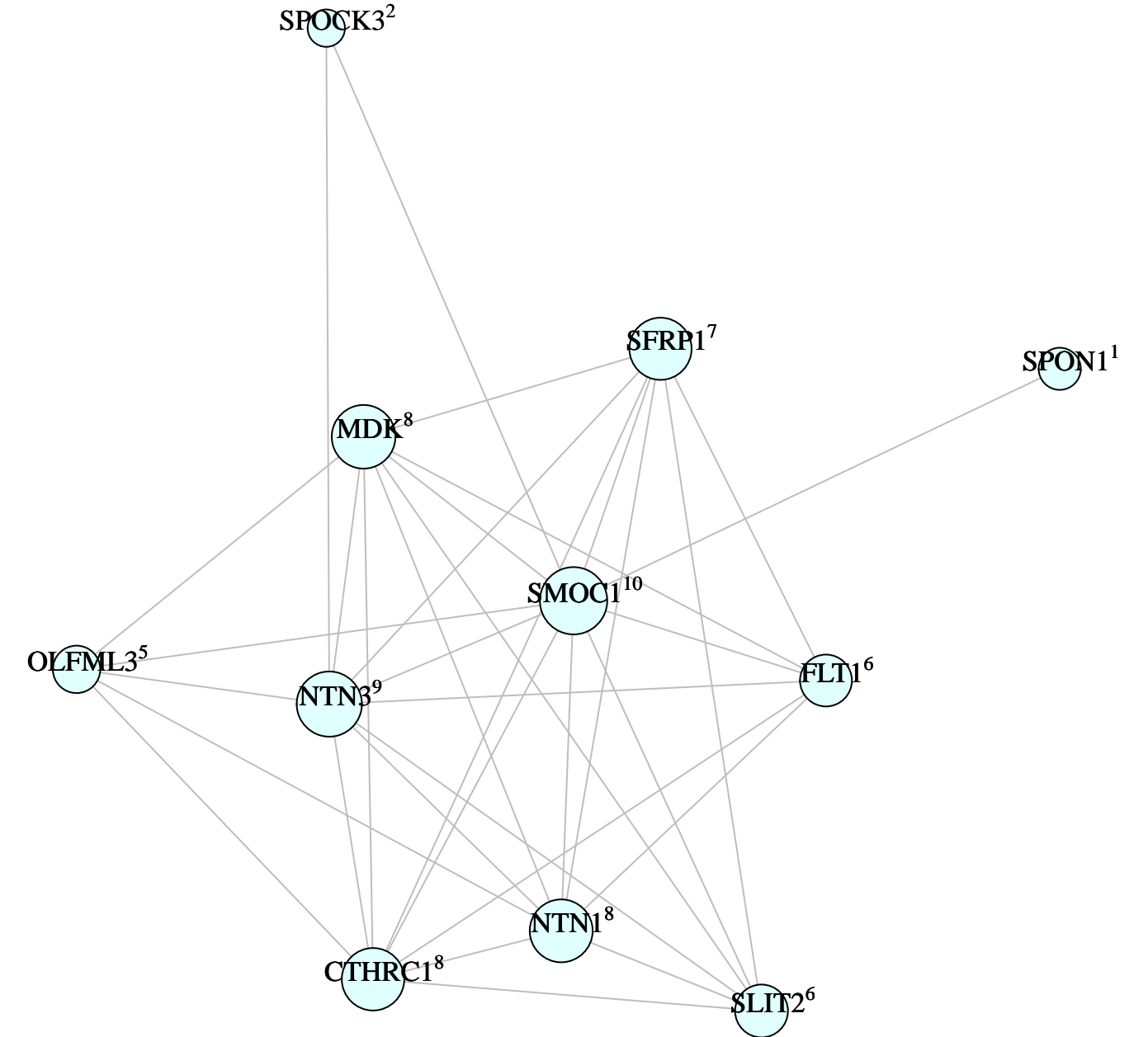
Network diagram showing interactions between various proteins. The central node is XRCC6, with a degree of 40. It is connected to many other nodes, including HNRNPM3, PRPF8, XPRC5, NCL, SSB, HNRNPR3, TRIM28, DDX23, ILF2, HDGF, RALY, HNRNUL1, NUDT21, RPA1, SF3A3, APEX1, HNRNPA3, KHDRBS3, UBA2, SAE1, SNRPD2, HNRNPH1, AMP32A, RBM25, HMGB1, SP100, HNRNPC3, SNRNP200, SNRNP70, THYN1, H2AFY1, HDGF1, and HDGF2. The nodes are represented by red circles, and the connections are gray lines.



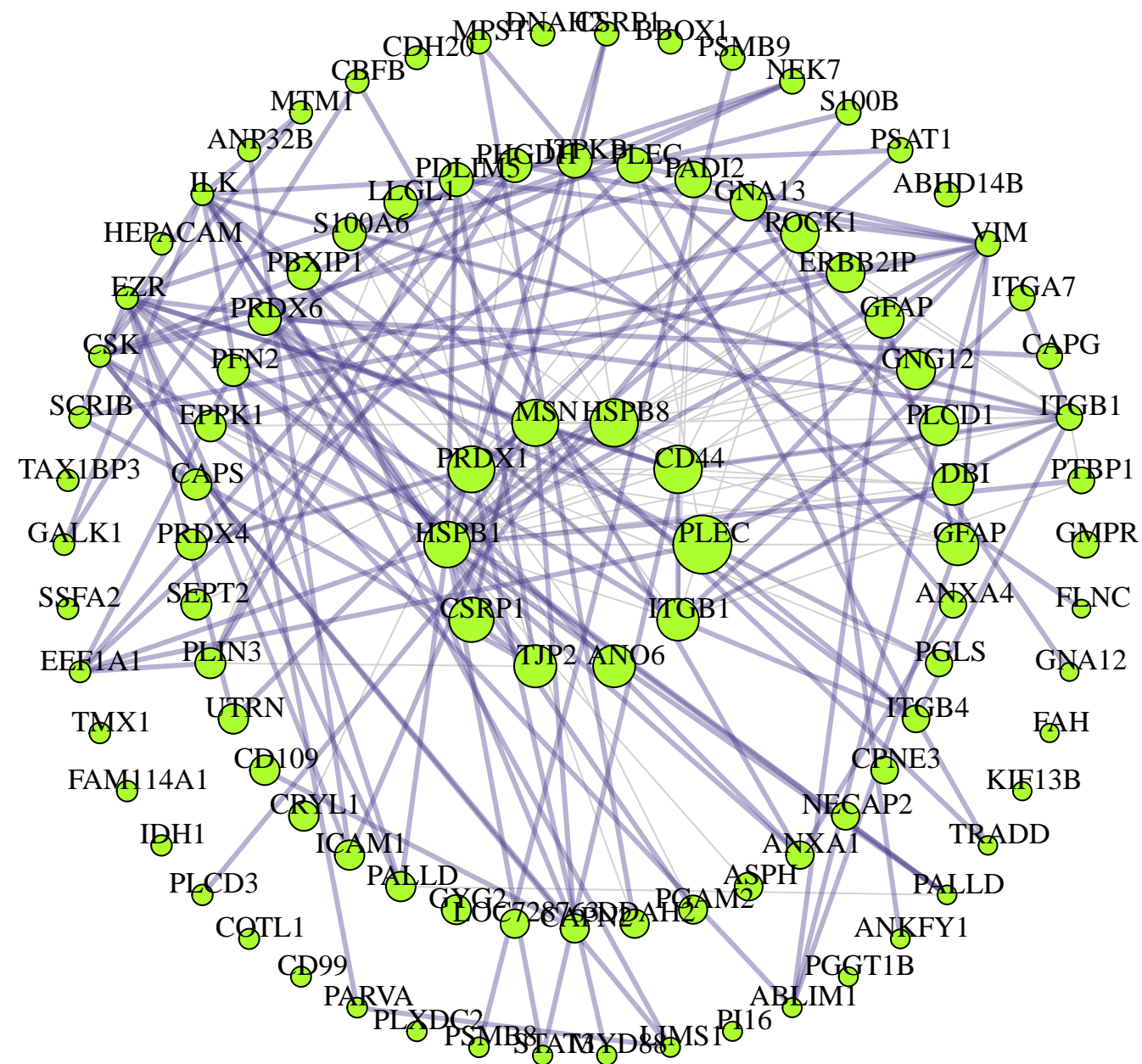
M42 lightcyan1 module



M42 lightcyan1 module hubs connected by top 70 TOM edges: HUB<sup>degree</sup>



## M11 greenyellow module



M11 greenyellow module hubs connected by top 150 TOM edges: HUB<sup>degree</sup>

