Using the shared genetics of dystonia and ataxia to unravel their pathogenesis

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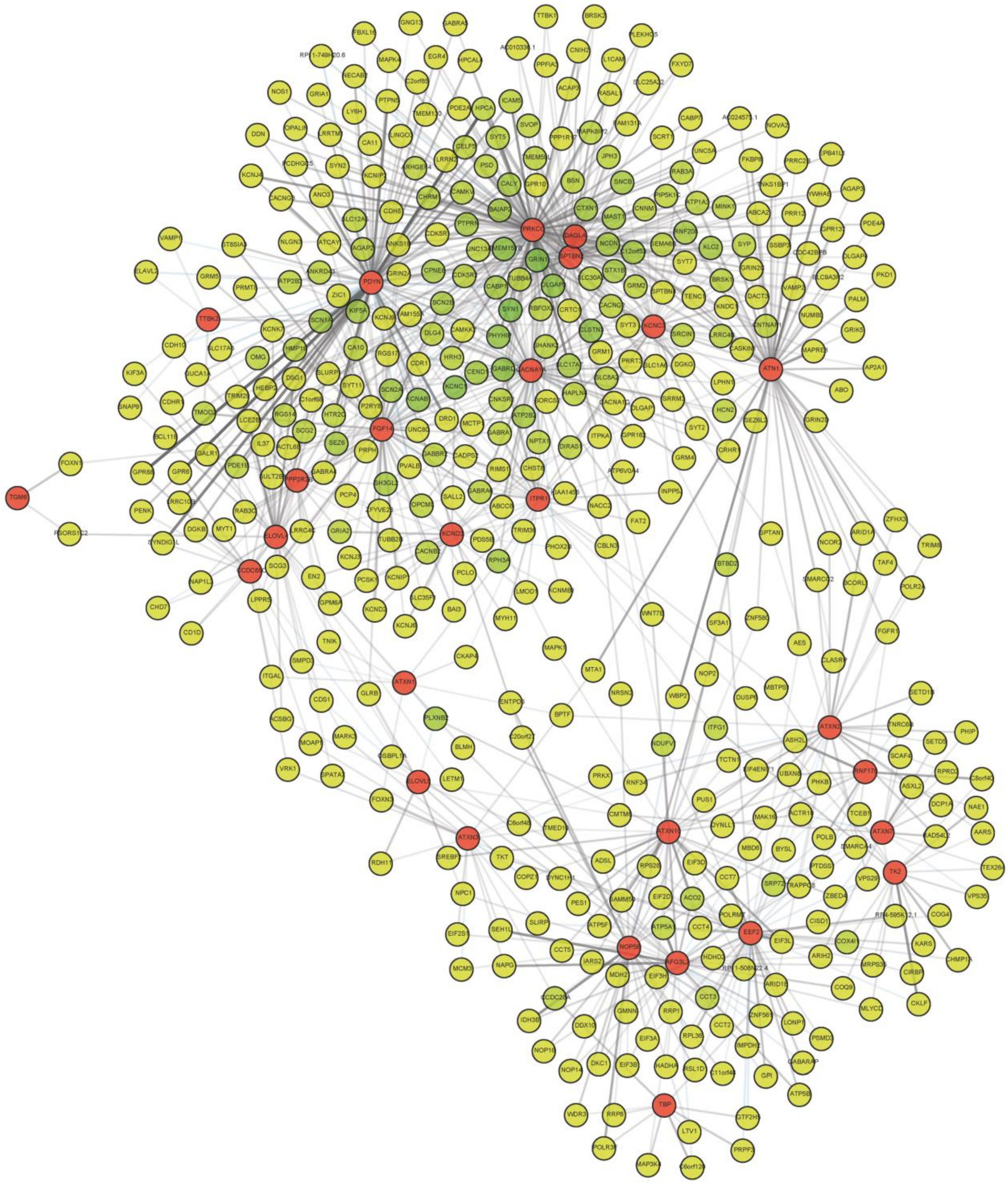
Supplementary Figure 1. Gene co-expression network based on known SCA genes. Red dots are the input genes (known SCA genes). Yellow to green dots represent co-expressed genes, darker colored dots have more connections. The thickness of the line indicates the strength of co-expression.



Supplementary Figure 2. Gene co-expression network based on known dystonia genes. Red dots are the input genes (known dystonia genes). Yellow to green dots represent co-expressed genes, darker colored dots have more connections. The thickness of the line indicates the strength of co-expression.