

The figure displays a network graph where nodes represent transcripts from specific tissues, and edges represent their relationships. The nodes are labeled as follows:

- transcript\_11-2\_ENSPTRG00000019474
- transcript\_3-19\_FBgn0010417
- transcript\_10-19\_ENSG00000106290
- transcript\_8-19\_ENSRNOG00000001355
- transcript\_10-20\_ENSG00000106290
- transcript\_7-19\_ENSMUSG000000036980
- transcript\_12-19\_ENSDARG00000102998
- transcript\_7-2\_ENSMUSG000000036980
- transcript\_3-46\_FBgn0010417
- transcript\_11-19\_ENSPTRG00000019474
- transcript\_3-40\_FBgn0010417
- transcript\_11-20\_ENSPTRG00000019474
- transcript\_10-2\_ENSG00000106290
- transcript\_8-2\_ENSRNOG00000001355
- transcript\_8-20\_ENSRNOG00000001355
- transcript\_3-20\_FBgn0010417
- transcript\_7-20\_ENSMUSG000000036980
- transcript\_12-20\_ENSDARG00000102998

The connections between these transcripts form a complex web, indicating shared or related functions across different biological contexts.