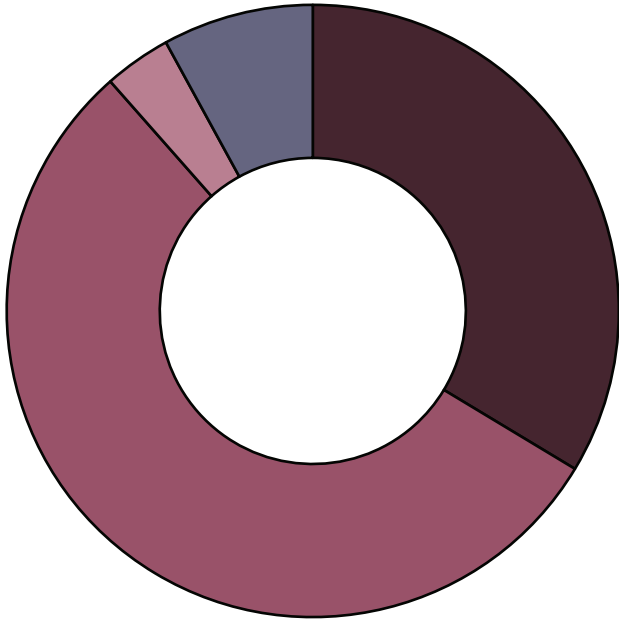
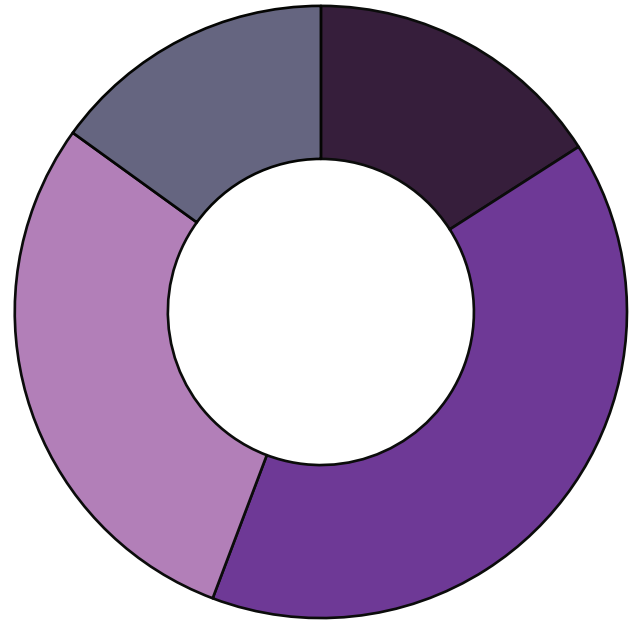


Function



Pathway



Negative	34%
Positive	55%
Variant	3%
Undetermined	8%

Extrinsic	16%
Intrinsic	40%
General	29%
Undetermined	15%

Apoptosis-related genes in *Euprymna scolopes*

Transcripts	293
Unique apoptosis genes	137