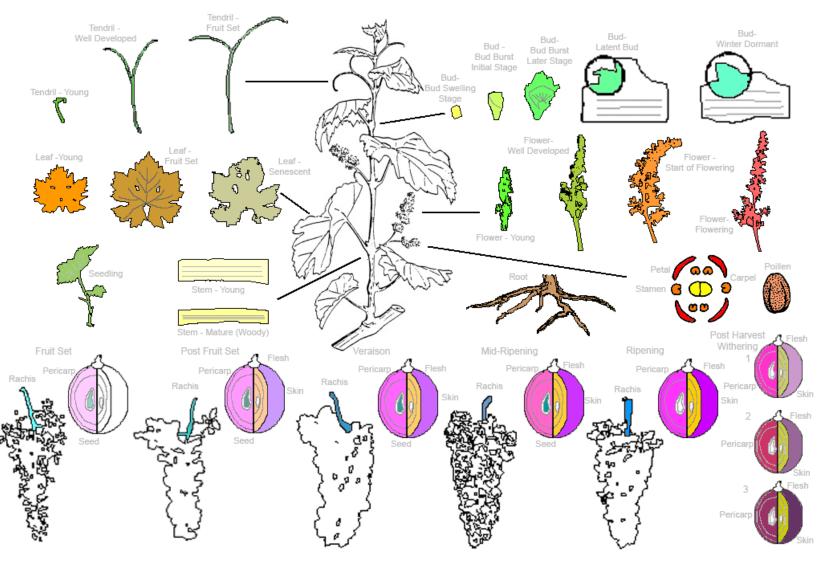
http://bar.utoronto.ca/efp_grape/cgibin/efpWeb.cgi

Data Source Mode □ Primary Gene ID Secondary Gene ID Signal Threshold □

? grape developmental ▼ Absolute ▼ VIT_00s0120g00060 VIT_09s0070g00080 500 Go

Grape eFP Browser at bar.utoronto.ca



eFP Browser by R. Patel. Images adapted and data derived from The Grapevine Expression Atlas Reveals a Deep Transcriptome Shift Driving the Entire Plant into a Maturation Program Marianna Fasoli et al., (2012) Plant Cell; 24(9): 3489–3505. Data were Nimblegen derived and were normalized using RMA and were provided as linearized data. All tissues were sampled in triplicate.

https://www.sdstate.edu/agronomy-horticulture-plant-science/function genomics-bud-endodormancy-induction-grapevines	<u>al-</u>

