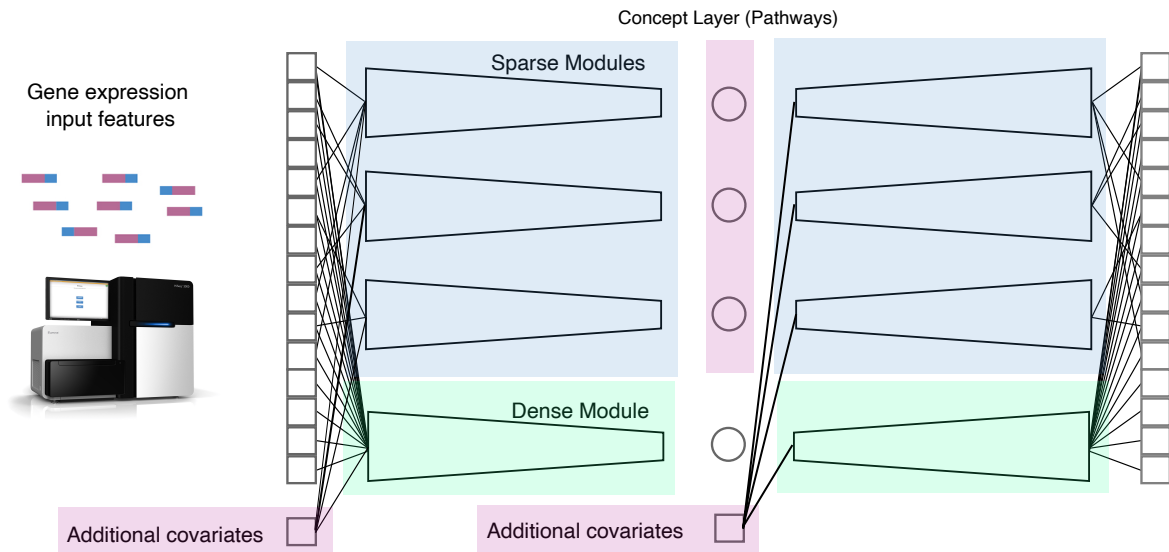
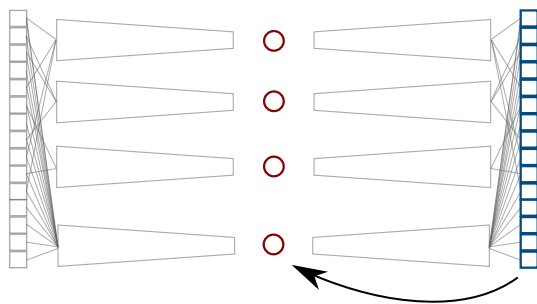


a**b***Attribute reconstruction loss to pathways***c***Attribute pathway outputs to input genes*