



Supplementary Figure S2: Local protein interaction network representing the significantly over-expressed druggable proteins for (a) Group 4, (b) Group 3 and (c) SHH MB. CIS-derived candidate cancer genes are represented by large circles, whilst small circles represent the proteins that interact with these CI-derived candidates. Nodes coloured in red-light red represent significantly over-expressed proteins, whilst blue-light blue represent significantly under-expressed proteins of that respective subgroup.