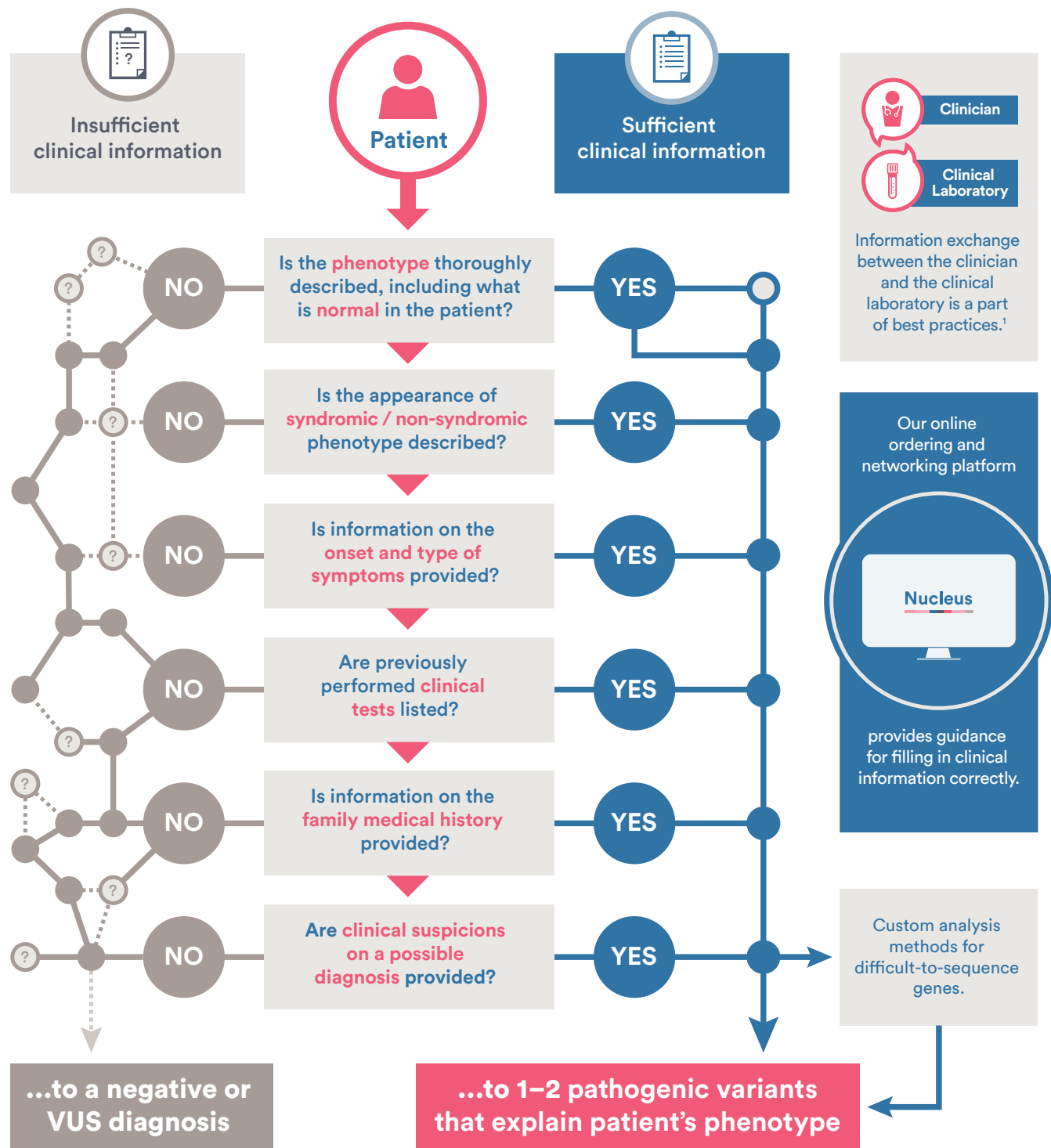


Clinical Information Can Lead to Finding a Variant That Might Otherwise Be Missed

From 20,000–35,000 possible variants explaining the phenotype...



¹ Bush et al. on behalf of the ACMG SELI committee. Genet Med. 2018 Feb; 20(2): 169–171.