

Supplementary Table S1. Overview of all *PLAG1*- and *HMGA2*-fusions so far described in pleomorphic salivary gland adenomas in the literature. The fusions were identified by RNA-sequencing, RT-PCR, fluorescence in situ hybridization (FISH), and/or array-based comparative genomic hybridization (arrayCGH).

Gene fusions	No. of cases	References
<i>PLAG1</i>-fusions		
<i>CTNNB1::PLAG1</i>	49	[1-9]
<i>CHCHD7::PLAG1</i>	19	[3,5,7,10]
<i>FGFR1::PLAG1</i>	10	[11]
<i>LIFR::PLAG1</i>	8	[3,5,12]
<i>TCEA1::PLAG1</i>	3	[2,10]
<i>FBXO32::PLAG1</i>	2	[7,13]
<i>NCALD::PLAG1</i>	3	[7]
<i>NTF3::PLAG1</i>	2	[7,13]
<i>ACTA2::PLAG1</i>	1	[7]
<i>DSTN::PLAG1</i>	1	[7]
<i>C1orf116::PLAG1</i>	1	[13]
<i>GEM::PLAG1</i>	1	[13]
<i>BOC::PLAG1</i>	1	[14]
<i>NFIB::PLAG1</i>	1	[6]
<i>HMGA2</i>-fusions		
<i>HMGA2::WIF1</i>	26	[3,5-9,15-17]
<i>HMGA2::NFIB</i>	5	[6,18]
<i>HMGA2::FHIT</i>	1	[19]
<i>HMGA2::RPSAP52</i>	1	[17]
<i>HMGA2::HELB</i>	1	[17]
<i>HMGA2::TMTC2</i>	1	[20]
<i>HMGA2::FTO</i>	1	[8]

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