

***NAB2-STAT6* fusion (Non-core)**

Reason/Evidentiary Support

In-frame *NAB2-STAT6* gene fusions result from chromosome 12q13 inversions and represent highly sensitive and specific signature alterations of meningeal solitary fibrous tumour/haemangiopericytoma (SFT/HPC) of grade 1, 2, or 3; these fusions are also characteristic of the analogous soft tissue/extracranial counterparts, which are referred to as SFT or malignant SFT. Given the relative ease of detecting this genetic alteration using a STAT6 immunohistochemical surrogate (see **STAT6 expression (immunohistochemistry)**), diagnostic confirmation is highly recommended in the WHO 2016 classification scheme before a diagnosis of SFT/HPC is rendered.^{1,2}

NAB2-STAT6 Gene Fusion

NAB2-STAT6 gene fusions are detectable using RT-PCR or various other sequencing techniques, including NGS if designed appropriately.^{3,4} Over 40 fusion variants have been detected to date, with the most common meningeal SFT/HPC subtypes fusing exon 6 of *NAB2* with exons 16, 17, or 18 of *STAT6* (roughly one-half of all cases).⁴ Preliminary data also suggests that the *NAB2* exon 4-*STAT6* exon 2/3 fusions are more common in the lower grade and clinically less aggressive SFT/HPC, though larger studies are needed for further validation.^{4,5}

STAT6 Nuclear Expression (Immunohistochemistry)

The STAT6 protein is normally expressed in the cytoplasm of cells, whereas *NAB2* is expressed in nuclei; however, the *NAB2-STAT6* fusions cause the STAT6 protein to translocate to the nucleus. As such, STAT6 immunohistochemistry represents a highly reliable and practical surrogate for detecting this signature alteration, with nearly 100% sensitivity and specificity regardless of the fusion variant.^{3,6} Nearly all meningeal SFT/HPC and extracranial SFTs display strong and extensive/diffuse nuclear positivity, whereas other diagnostic considerations, such as meningiomas, nerve sheath tumours, and various sarcomas, either lack expression or show only cytoplasmic staining. As such, the pathologist is cautioned against rendering a diagnosis of SFT/HPC in the absence of nuclear STAT6 immunoreactivity.

References

- 1 Louis DN, Perry A, Reifenberger G, von Deimling A, Figarella-Branger D, Cavenee WK, Ohgaki H, Wiestler OD, Kleihues P and Ellison DW (2016). The 2016 World Health Organization Classification of Tumors of the Central Nervous System: a summary. *Acta Neuropathologica* 131(6):803-820.
- 2 Louis DN, Ohgaki H, Wiestler OD and Cavenee WK (eds) (2016). *WHO Classification of Tumours of the Central Nervous System, Revised. Fourth Edition*, IARC, Lyon.
- 3 Schweizer L, Koelsche C, Sahm F, Piro RM, Capper D, Reuss DE, Pusch S, Habel A, Meyer J, Gock T, Jones DT, Mawrin C, Schittenhelm J, Becker A, Heim S, Simon M, Herold-Mende C, Mechttersheimer G, Paulus W, Konig R, Wiestler OD, Pfister SM and von Deimling A (2013). Meningeal hemangiopericytoma and solitary fibrous tumors carry the *NAB2-STAT6* fusion and can be diagnosed by nuclear expression of STAT6 protein. *Acta Neuropathol* 125(5):651-658.

- 4 Nakada S, Minato H and Nojima T (2016). Clinicopathological differences between variants of the NAB2-STAT6 fusion gene in solitary fibrous tumors of the meninges and extra-central nervous system. *Brain Tumor Pathol* 33(3):169-174.

- 5 Fritchie KJ, Jin L, Rubin BP, Burger PC, Jenkins SM, Barthelmess S, Moskalev EA, Haller F, Oliveira AM and Giannini C (2016). NAB2-STAT6 Gene Fusion in Meningeal Hemangiopericytoma and Solitary Fibrous Tumor. *J Neuropathol Exp Neurol* 75(3):263-271.

- 6 Koelsche C, Schweizer L, Renner M, Warth A, Jones DT, Sahm F, Reuss DE, Capper D, Knosel T, Schulz B, Petersen I, Ulrich A, Renker EK, Lehner B, Pfister SM, Schirmacher P, von Deimling A and Mechttersheimer G (2014). Nuclear relocation of STAT6 reliably predicts NAB2-STAT6 fusion for the diagnosis of solitary fibrous tumour. *Histopathology* 65(5):613-622.