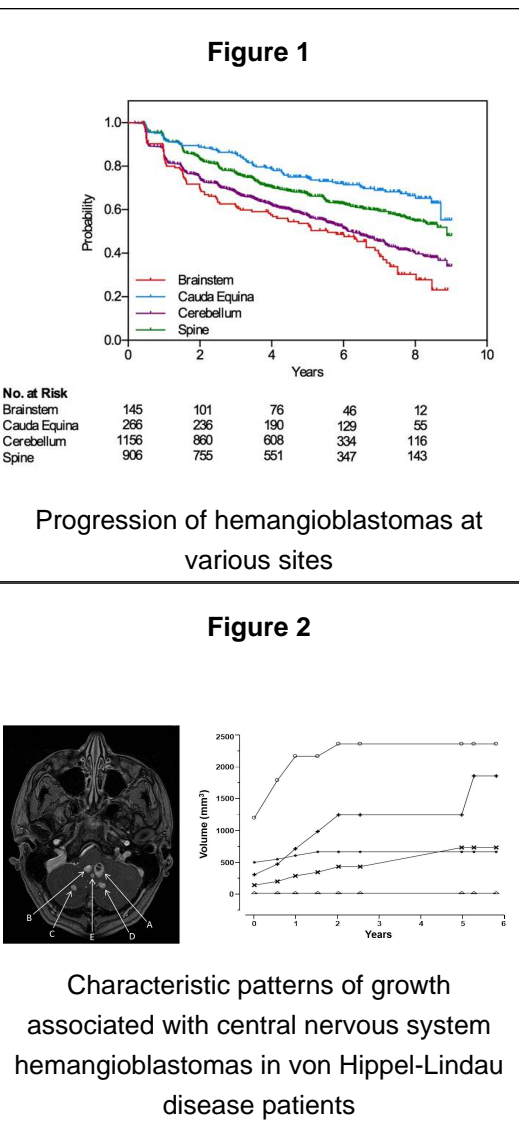


By the conclusion of this presentation, participants should be able to: 1) describe the natural history of central nervous system hemangioblastomas, 2) describe the factors that impact tumor development, growth and symptom formation and 3) identify most effective treatment strategies for these neoplasms.

Increased tumor burden was associated with partial deletions in VHL gene ( $P=0.005$ ) and male sex ( $P=0.002$ ). Hemangioblastoma development (median, 0.3 new tumors/year) was associated with younger age ( $P<0.0001$ ) and more tumors at study entrance ( $P<0.0001$ ). While 1,278 hemangioblastomas (51%) did not grow, 1,226 hemangioblastomas (49%) grew in a saltatory (886 tumors; 72% of growing tumors), linear (76; 6%), or exponential (264; 22%) pattern. Faster tumor growth was associated with male sex ( $P=0.002$ ), symptomatic tumors ( $P<0.0001$ ) and tumors associated with cysts ( $P<0.0001$ ). Location-dependent tumor size was the primary predictor of eventual symptom formation (159 symptomatic tumors [6.4%]; area under the curve greater than 0.9).



CNS hemangioblastoma burden in VHL is associated with partial germline deletions and male sex. Unpredictable growth of hemangioblastomas compromises assessment of non-surgical therapies. Judicious treatment of symptom-producing hemangioblastomas, while avoiding unnecessary treatment of asymptomatic tumors that may not progress, can provide clinical stability.

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