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Progression in patients with low- and intermediate1-risk del(5q)
myelodysplastic syndromes is predicted by a limited subset of
mutations

Christian Scharenberg, Valentina Giai, Andrea Pellaatti, Leonie Saft,
Marios Dimitriou, Monica Jansson, Martin Jädersten, Alf Grandien,
Iyadh Douagi, Donna S. Neuberg, Katarina LeBlanc, Jacqueline
Boultwood, Mohsen Karimi, Sten Eirik W. Jacobsen, Petter S. Woll, and
Eva Hellström-Lindberg

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