

Durvalumab plus gemcitabine and cisplatin in patients with advanced biliary tract cancer: an exploratory analysis of real-world data

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Supplementary Table 1: Clinically relevant genetic alterations

Patient	TMB	gene	type of mutation
2	4.72	<i>IDH1</i>	activating mutation
2	4.72	<i>BAP1</i>	deletion mutation
8	3.91	<i>MSH6</i>	deletion mutation
8	3.91	<i>GLI1</i>	deletion mutation
8	3.91	<i>BAP1</i>	deletion mutation
9	4.71	<i>TP53</i>	deletion mutation
9	4.71	<i>MUTYH</i>	deletion mutation
9	4.71	<i>ARID2</i>	deletion mutation
12	n.a.	<i>ERBB3</i>	activating mutation
12	n.a.	<i>BRAF</i>	deletion mutation
13	5.49	<i>TP53</i>	deletion mutation
13	5.49	<i>ABL1</i>	deletion mutation
13	5.49	<i>RTN3::EWSR1</i>	fusion
14	10.16	<i>NRAS</i>	deletion mutation
14	10.16	<i>TP53</i>	deletion mutation
14	10.16	<i>SMAD4</i>	deletion mutation
14	10.16	<i>AR</i>	deletion mutation
15	3.13	<i>KRAS</i>	activating mutation
16	7.03	<i>KRAS</i>	activating mutation
16	7.03	<i>SMAD4</i>	deletion mutation
18	3.91	<i>ARAF</i>	activating mutation
18	3.91	<i>BAP1</i>	deletion mutation
18	3.91	<i>LRP1B</i>	deletion mutation
20	3.91	<i>BAP1</i>	deletion mutation
20	3.91	<i>RANBP2</i>	deletion mutation
20	3.91	<i>FGFR2::NEK9</i>	fusion
34	0.00	<i>RET</i>	deletion mutation
34	0.00	<i>FGFR2::KCNH7</i>	fusion