

No Benefit of BCR-ABL1 Screening in Polycythemia

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Dear Sir,

Chronic Myeloid Leukaemia (CML) has become the paradigm for the rational development of a molecularly targeted therapy in oncology and is characterised morphologically by leukocytosis due to neutrophils in different stages of maturation, no significant dysplasia, low blast cell numbers, a variable monocytosis, commonly basophilia and/or eosinophilia, and a variable platelet count. The genetic hallmark of CML is (9;22)(q34;q11.2) resulting in the BCR-ABL1 fusion gene [1]. Erythrocytosis is an increase in the number of red blood cells and is a frequent reason for haematological investigation. Common causes include hypoxia (smoking, lung or cardiac disease, sleep apnoea), drugs (diuretics, testosterone, or anabolic steroids, erythropoietin) or the myeloproliferative neoplasm polycythemia vera (PV) [2]. PV is characterised at the molecular level by the presence of JAK2 exon 14 V617F and exon 12 insertion or deletion mutations [3]. The term polycythemia is often used synonymously with erythrocytosis however this is incorrect as polycythemia implies a leukocytosis and thrombocytosis in addition to the increase in red cells. As both CML and PV can present with leukocytosis and thrombocytosis, testing for BCR-ABL1 can often be erroneously triggered in such cases regardless of the presence of an erythrocytosis.

In order to determine whether any continued value exists in screening for the BCR-ABL1 fusion gene as part of the investigation in patients with polycythemia and erythrocytosis, a retrospective audit was performed. In a ten year period from January 2006 to December 2015 inclusive, 6578 diagnostic requests were received for BCR-ABL1 transcript identification at a national testing centre for haematological malignancies. Detection of BCR-ABL1 transcripts

was performed according to standardized protocols and guidelines. Of these 6578 requests, 36 (0.5%) had clinical details stated as raised haemoglobin and haematocrit (n=17), polycythemia (n=16) and erythrocytosis (n=3). BCR-ABL1 transcripts were not detected in any of these 36 patients.

The absence of BCR-ABL1 transcripts in patients with polycythemia or erythrocytosis suggests that routine exclusion of CML by BCR-ABL1 fusion transcript identification is not warranted in such cases. However, extremely rare confounding scenarios should always be considered and include the co-existence of CML and PV and singular cases of CML presenting with polycythemia [4,5]. While JAK2 mutation analysis for PV remains high in the diagnostic algorithm for investigation of polycythemia and erythrocytosis [2], more common reasons should be excluded before subsequent investigation of rare underlying causes.

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