



# Online Individual Assessment for classification of BRCA1 and BRCA2 variants 2021 (Run 6)

Complete the assessment: 29th April – 31st May 2021

The aim of this module is to provide an online competency assessment for the classification of **BRCA1** and **BRCA2** variants. This module is supported by an educational grant from AstraZeneca and is delivered jointly by EMQN CIC and GenQA.



This resource is delivered using the **GenQA Genomics Training** and **Competence Assessment Tool (GTACT)**. GTACT is webbased and can be accessed from any global location, therefore there are no limits to the number of participants.

This assessment is open to individuals who classify the pathogenicity of variants and provides assurance of their competency to achieve this accurately and in the most up-to-date manner.

## Information on participation

- There is no fee to participate
- Once registered for Run 6, all participants will receive:
  - > full instructions for Run 6 via email
  - registration details for two educational webinars.

#### Webinar 1

**BRCA1** and **BRCA2** variant classification masterclass

28th April 2021 1-2pm BST

#### Webinar 2

Discussion of results from Run 6 assessment

17th June 2021 1-2pm BST

- Secure website access to GTACT will be provided.
- Participants submit their classification and evidence for the six variants.
- Please use **ACMG Variant Classification guidelines** (Richards *et al.*, 2015)

### **Benefits**

All participants will receive the following:

- Individual report summarising their classification,
- **Summary report** to enable benchmarking against the expert predicted classifications and details of the evidence used,
- Certificate of participation.

Your participation in this assessment and your individual data, remain confidential between yourself and EMQN / GenQA.

For further details and to register please go to <a href="https://genga.org/BRCA-HRR">https://genga.org/BRCA-HRR</a>

Note: Run 7 will be provided in November 2021 and will focus on variants in other HRR genes.