## COELIAC UK RESEARCH CONFERENCE 2017 CPD QUESTIONS



## Decoding the genetic background of coeliac disease – Dr Iris Jonkers

Please type your answers below and print for your records or print this document and complete by hand.

| Q1 | What is a genome wide association study (GWAS)?   |
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| А  |   |
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| Q2 | Which genes are known to be needed for a person to develop coeliac disease?   |
| A  | Which genes are known to be neceded for a person to develop coeffic disease.  |
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| Q3 | Loci associated with coeliac disease are also associated with other autoimmune diseases. What are three examples of other autoimmune diseases with loci which |
|    | overlap with coeliac disease?   |
| А  |   |
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| Q4 | More than 95% of single nucleotide polymorphisms (SNPs) associated with   |
|    | coeliac disease fall in non-coding regions, how can the role of these SNPs in   |
| Δ. | coeliac disease be explained?   |
| Α  |   |
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| Q5 | Which genes are thought to lead to altered barrier function in coeliac disease?   |
| А  |   |
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After answering these questions, should you require any further information, please contact us at <a href="mailto:ruth.passmore@coeliac.org.uk">ruth.passmore@coeliac.org.uk</a>. You can also discuss this further on our HCP forum: <a href="https://www.coeliac.org.uk/hcp-forum">www.coeliac.org.uk/hcp-forum</a>.