<u>Immunity – AS 9700 Biology Nov 2022</u>

1. Nov/2022/Paper 11/No.40

A student wrote the following statements about the production of monoclonal antibodies.

Which statement is not correct?

- A Non-self antigen is injected into a mammal.
- **B** Plasma cells secrete the specific antibody in response to a non-self antigen.
- C Selected hybridoma cells divide by mitosis to form a clone.
- **D** T-lymphocytes are fused with cancer cells to form hybridoma cells.

2. Nov/2022/Paper_12/No.35

Which statement about the structure of the heart is correct?

- A The semilunar valves close when the pressure in the ventricles falls lower than the pressure in the arteries.
- **B** A muscle called the septum separates the atria from the ventricles.
- C There is a small patch of tissue in the right atrium that acts to delay the electrical impulse.
- **D** Closure of valves in the veins prevents backflow of blood into the ventricles.

3. Nov/2022/Paper 12/No.39

Immunity to certain pathogens develops when a person is given a vaccination.

Which effects of vaccination are correct?

- 1 production of antibodies to protect against future infections
- 2 results in artificial active immunity
- 3 stimulation of appropriate lymphocytes
- **A** 1, 2 and 3 **B** 1 and 2 only **C** 1 and 3 only **D** 2 and 3 only

4. Nov/2022/Paper 12/No.40

A student wrote three statements about antibodies.

- 1 Their structure depends on peptide, hydrogen and disulfide bonds.
- 2 They are protein molecules with both tertiary and quaternary structure.
- 3 Four polypeptides provide four antigen binding sites.

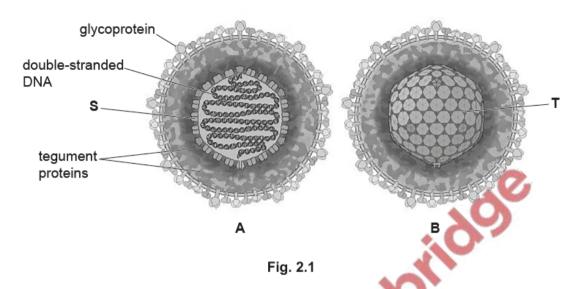
Which statements are correct?

A 1, 2 and 3 **B** 1 and 2 only **C** 1 and 3 only **D** 2 and 3 only

5. Nov/2022/Paper_22/No.2(a)

Human cytomegalovirus (HCMV) is a common virus affecting humans. In people with a fully functioning immune system, infection by HCMV usually causes no, or only mild, symptoms.

Fig. 2.1A is a diagram of a section through HCMV. In Fig. 2.1B, only the outer part of HCMV is sectioned.



The viral DNA shown in Fig. 2.1 contains genes that code for proteins important in viral replication and viral structure, including viral DNA polymerase and proteins known as tegument proteins.

Viruses can only replicate in host cells as they need to use processes and contents of the host cell. Complete viral particles that are released from the host cell are known as virions.

(a) Structure S in Fig. 2.1A is a subunit of structure T in Fig. 2.1B.

Name the chemical compound used to make structure S and name	structure I.	•
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S		N
т	X	
	•	[2]

mala	aria.	dium falciparum is one species of <i>Plasmodium</i> that causes the life-threatening disease With early diagnosis and the correct drug treatment, the pathogen can be eliminated from a particularly if the disease is not severe.								
(a)	Nan	ne the type of pathogen that causes malaria.								
		[1]								
(b)	To help prevent the development and spread of drug resistance in <i>Plasmodium</i> , the Health Organization (WHO) recommends using a treatment known as artemisinin-combination therapy (ACT).									
	AC	Γ involves two different types of drug:								
		a fast-acting drug derived from a compound known as artemisinin, which causes a rapid decrease in the number of <i>P. falciparum</i> one or more longer-acting, non-artemisinin, drugs that eliminate any remaining pathogens.								
	(i)	Suggest why using ACT with the two different types of drug is more effective in preventing the development of drug resistance in <i>Plasmodium</i> than a treatment using only one type of drug.								
		[2]								
	(ii)	In some areas, partial artemisinin resistance has developed. This means ACT takes a longer time for the pathogen to be eliminated from the body.								
		Explain why there is an increased risk of transmission of the pathogen to other people if a person is receiving ACT and the pathogen has partial artemisinin resistance.								
		[2]								

6. Nov/2022/Paper_22/No.3(a_ b)

7. Nov/2022/Paper_23/No.2(a)

Fig. 2.1 is a scanning electron micrograph showing a macrophage engulfing some bacteria.

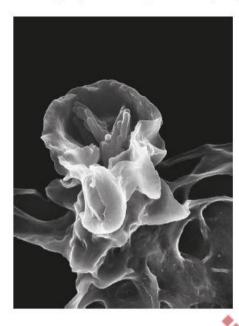


Fig. 2.1

(a)	(i)	Describe how macrophages engulf bacteria.							

Palpacamilla idde

(ii)

8.	Cath One	elici role	22/Paper_23/No.5(a_b) licidin LL-37 is a cell signalling compound that stimulates many different cells in humans. ole of cathelicidin LL-37 is stimulating the production of endothelial cells in the formation of the during wound healing.											
	(a)	(i)		n how i ing com		sible fo	r many	differen	t cell ty	pes to	respond	to the	same o	ell
					•••••									
												•••••		
					•••••					•••••		•••••		
														[2]
	((ii)	Descri	be the a	ppearar	nce of th	e endot	helial ce	ells of a	capillary	7			
												•••••		
									P	*		•••••		
				••••••								•••••		
					•••••			,						[2]
						A.	0							[—]
Cathelicidin LL-37 is a protein composed of 37 amino acids. Table 5.1 shows:														
			40	sequenc					-					
				IA triplets the prima					e gene th	at codes	for the fi	rst 10 am	ino acids	į
							Ta	able 5.1						
		6	mino acid osition	1	2	3	4	5	6	7	8	9	10	

asp

GAT

phe

TTC

phe

TTC

arg

CGG

lys

AAA

ser

TCT

lys

AAA

amino

acid DNA

triplet

leu

CTG

leu

CTG

gly

GGT

Table 5.2

	second base												
			Т		С		Α		G				
			TTT		TCT		TAT	tur	TGT	CVC	Т		
		т	TTC	phe	TCC	ser	TAC	tyr	TGC	cys	С		
		'	TTA	leu	TCA	361	TAA	stop	TGA	stop	Α		
			TTG	ieu	TCG		TAG	stop	TGG	try	G		
			CTT		CCT		CAT	his gln	CGT		Т		ψ
_ n		С	CTC	leu	CCC	pro	CAC		CGC	arg	С		
base		_	CTA	ieu	CCA		CAA		CGA	u g	Α		third base
first b			CTG		CCG		CAG	giii	CGG		G		힏
Įį.		Α	ATT	ile	ACT	thr	AAT	asp	AGT	ser	1		‡.
			ATC		ACC		AAC	aop	AGC		С		
		^	ATA	ile	ACA		AAA	lys	AGA	arg	Α		
			ATG	met	ACG		AAG	.,,	AGG	9	G		
			GTT		GCT		GAT	asp	GGT		Т		
		G	GTC	val	GCC	ala	GAC		GGC	gly	С		
		٦	GTA		GCA		GAA	glu	GGA	9'7	Α		
			GTG		GCG		GAG	and .	GGG		G		

(b)	Mutations of DNA base sequences	in a	gene	can affect t	the primary	structure of	proteins
٠,		-	-				

Use the information in Table 5.1 and Table 5.2 to suggest the effect on the primary structure of cathelicidin LL-37 of:

(i)	the substitution of the base T with the base A in the middle of the triplet at position 5	[1]
(ii)	the deletion of base T in the triplet at position 2	
		[2]
(iii)	the insertion of base G between bases G and T in the triplet at position 3.	