

7.9 Phylogeny

Name: _____ Class: _____ Date: _____

Total: 9 marks**KEY WORDS** phylogenetic tree 系统发生树 • evolution 进化 • common ancestor 共同祖先

Objective

Build the skills to answer exam questions on **phylogeny** — reading evolutionary trees.**You must be able to:**

- read a **phylogenetic tree** 系统发生树 (branches, nodes, common ancestors)
- identify the **most recent common ancestor** of two species
- interpret shared derived traits

1 Worked examples

Study these first. Each one shows the method for a question type used later — follow the steps and you can do the Practice and Exam-style questions yourself.

■ What a tree shows

A **phylogenetic tree** shows how species are related by descent. **Tips** are species; **nodes** (branch points) are **common ancestors**; the closer two tips branch, the more closely related.

■ Most recent common ancestor

The **most recent common ancestor** of two species is the node where their lines last join — trace each back until the branches meet.

■ Reading relatedness

Two species that share a **more recent** node are **more closely related** than either is to a species that branches off earlier.

■ Shared derived traits

A trait shared by all descendants of a node (and not by outgroups) marks that branch — used to build and read the tree.

2 Practice

Now apply the methods above.

2.1 What does a node (branch point) on a tree represent? [1]**2.2** How do you find the most recent common ancestor of two species? [1]**2.3** Are two species that branch off close together more or less related? [1]

3 Exam-style questions

3.1 On a phylogenetic tree, the tips represent [1]

A	B
C	D

A common ancestors**B** species (or groups)**C** mutations**D** fossils only**3.2** On a tree, species A and B branch from one node; species C branches off earlier.

(a) Which two species are most closely related? [1]

(b) Explain your reasoning using the tree. [2]

3.3 Explain how shared derived traits are used to group species on a tree. [2]