Finding Normality in Abnormality: Normal Function Ascription in Cancer Biology

Seth Goldwasser University of Pittsburgh

Outline 1. Normal Function and Function Pluralism

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To a first approximation, normal functions are a 4-place predicate.

An activity, *φ*, of a part, *p*, of a token system, *s*, is a <u>normal function</u>, F, iff items of p's type make contributions, *C*, to systems of s's type by *φ*-ing (cf. Weber 2017).

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- That said, different types of function are ascribed in different sub-disciplines of biology.
- Causal role or "minimal" functions are ascribed in cladistic systematics, while normal functions are ascribed in physiology.
- If an account of function aims to explicate the ascription of function in a subdiscipline then it aims to be descriptive.
- There are at least two desiderata on descriptive accounts of function



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Contribution



<u>Class Adequacy</u>: a descriptive account should be extensionally adequate concerning the types of system to which a sub-discipline of biology ascribes a type of function.



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•Example: assuming ecosystems don't have parts with normal functions (narrow).

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Accounts can be criticized for entailing conditions on the ascription of function to which biologists do not adhere.

•Example: Amundson and Lauder (1994) and Griffiths (1994, 2006) claim that cladists do not nor need to appeal to history when ascribing a function.

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The standard classificatory scheme is helpful <u>but not</u> <u>perfect</u>.

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A Representative Example: Peinado et al. 2012

Peinado et al. (2012) features the ascription of a normal function to part of melanoma.

Increased MET levels... and MET activation in bone marrow cells

Peinado and Colleagues Ascribe a Normal Function

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The experiments they ran were designed to figure out how sEV contribute to premetastatic niche construction.

The ascription tells us how sEV normally so contribute. That is, the function ascription identifies a <u>standard</u> for sEV activity, disposition, and structure.

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Selected effects accounts claim that an activity of a part is a normal functions if and only if that part <u>is **selected**</u> <u>for performing that activity</u>.

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For a trait to be subject to selection, the system(s) in which that trait is present has to meet at least three conditions:

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1.The system(s) has to exhibit *heritable variants* of the trait or trait-type.

2. The **fit** between system(s) and environment has to **favor** <u>some traits over others</u>.

3. The selected trait has to be **retained over variants**.

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At the cellular level, parts of cancers that are ascribed normal functions are often the product of drift or genetic hitchhiking without being co-opted.

At the tumor level, these parts are often neither heritable or recapitulated in metastases nor the product of competition between tumors.

In which case, at least some parts of cancers have normal functions despite not being subject to selection

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Another type of normality that's thought to be scientifically respectable is *conditional statistical typical ity*.

The normal function of a part then becomes what that part does to contribute to the (inclusive) fitness of an individual system which, conditional on that part's so contributing, is typical for systems of the type.

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But doing so threatens to undermine the normal function (pun intended) of ascribing normal functions, namely, identifying stan dards of the activities, dispositions, and structural features of biological systems.

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An Objection

Moreover, cancer biologists reserve talk of "normal functioning" to describe the healthy variants of the disease processes they study. In response, if we understand normal functioning as a part's activity embodying a standard then cancer biologists are engaged in the practice of ascribing those functions.

A Reply

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The reason for the practice is clinical: if there are normal ways that a disease progresses then understanding that standard aids undermining it.

A Reply

In response, if we understand normal functioning as a part's activity embodying a standard then cancer biologists are, in fact, engaged in the practice of ascribing those functions.

The reason for the practice is clinical: if there are normal ways that a disease progresses then understanding that standard aids undermining it.

The ascription of these functions is helpful here because cancers are so variable.

A Reply

Summary

1. Different types of function are ascribed in across biology.

2. Descriptive accounts of function must satisfy class adequacy and explanatory adequacy.

3. Cancer biology features the ascription of normal *(abnormality inclusive functions.)*

4. Selected-effects Accounts and Fitness Contribution Accounts struggle to account for this.

