# WikipediA Singular homology

In <u>algebraic topology</u>, **singular homology** refers to the study of a certain set of <u>algebraic invariants</u> of a <u>topological space</u> *X*, the so-called **homology groups**  $H_n(X)$ . Intuitively, singular homology counts, for each dimension *n*, the *n*-dimensional holes of a space. Singular homology is a particular example of a <u>homology</u> <u>theory</u>, which has now grown to be a rather broad collection of theories. Of the various theories, it is perhaps one of the simpler ones to understand, being built on fairly concrete constructions (see also the related theory simplicial homology).

In brief, singular homology is constructed by taking maps of the <u>standard *n*-simplex</u> to a topological space, and composing them into <u>formal sums</u>, called **singular chains**. The boundary operation – mapping each *n*-dimensional simplex to its (n-1)-dimensional <u>boundary</u> – induces the singular <u>chain complex</u>. The singular homology is then the <u>homology</u> of the chain complex. The resulting homology groups are the same for all <u>homotopy equivalent</u> spaces, which is the reason for their study. These constructions can be applied to all topological spaces, and so singular homology is expressible as a <u>functor</u> from the <u>category of topological spaces</u> to the category of graded abelian groups.

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### **Singular simplices**

A <u>singular *n*-simplex</u> in a topological space *X* is a <u>continuous function</u> (also called a map)  $\sigma$  from the standard *n*-<u>simplex</u>  $\Delta^n$  to *X*, written  $\sigma : \Delta^n \to X$ . This map need not be <u>injective</u>, and there can be non-equivalent singular simplices with the same image in *X*.

The boundary of  $\sigma$ , denoted as  $\partial_n \sigma$ , is defined to be the <u>formal sum</u> of the singular (n - 1)-simplices represented by the restriction of  $\sigma$  to the faces of the standard *n*-simplex, with an alternating sign to take orientation into account. (A formal sum is an element of the <u>free abelian group</u> on the simplices. The basis for the group is the infinite set of all possible singular simplices. The group operation is "addition" and the sum of simplex *a* with simplex *b* is usually simply designated *a* + *b*, but a + a = 2a and so on. Every simplex *a* has a negative -a.) Thus, if we designate  $\sigma$  by its vertices

 $[p_0,p_1,\ldots,p_n]=[\sigma(e_0),\sigma(e_1),\ldots,\sigma(e_n)]$ 

corresponding to the vertices  $e_k$  of the standard *n*-simplex  $\Delta^n$  (which of course does not fully specify the singular simplex produced by  $\sigma$ ), then



The standard 2-simplex  $\Delta^2$  in  $\mathbf{R}^3$ 

$$\partial_n \sigma = \partial_n [p_0, p_1, \dots, p_n] = \sum_{k=0}^n (-1)^k [p_0, \dots, p_{k-1}, p_{k+1}, \dots, p_n] = \sum_{k=0}^n (-1)^k \sigma|_{e_0, \dots, e_{k-1}, e_{k+1}, \dots, e_n}$$

is a <u>formal sum</u> of the faces of the simplex image designated in a specific way.<sup>[1]</sup> (That is, a particular face has to be the restriction of  $\sigma$  to a face of  $\Delta^n$  which depends on the order that its vertices are listed.) Thus, for example, the boundary of  $\sigma = [p_0, p_1]$  (a curve going from  $p_0$  to  $p_1$ ) is the formal sum (or "formal difference")  $[p_1] - [p_0]$ .

### Singular chain complex

The usual construction of singular homology proceeds by defining formal sums of simplices, which may be understood to be elements of a <u>free abelian group</u>, and then showing that we can define a certain group, the **homology group** of the topological space, involving the boundary operator.

Consider first the set of all possible singular *n*-simplices  $\sigma_n(X)$  on a topological space *X*. This set may be used as the basis of a free abelian group, so that each singular *n*-simplex is a generator of the group. This set of generators is of course usually infinite, frequently <u>uncountable</u>, as there are many ways of mapping a simplex into a typical topological space. The free abelian group generated by this basis is commonly denoted as  $C_n(X)$ . Elements of  $C_n(X)$  are called **singular** *n***-chains**; they are formal sums of singular simplices with integer coefficients.

The <u>boundary</u>  $\partial$  is readily extended to act on singular *n*-chains. The extension, called the <u>boundary operator</u>, written as

 $\partial_n: C_n o C_{n-1},$ 

is a <u>homomorphism</u> of groups. The boundary operator, together with the  $C_n$ , form a <u>chain complex</u> of abelian groups, called the **singular complex**. It is often denoted as  $(C_{\bullet}(X), \partial_{\bullet})$  or more simply  $C_{\bullet}(X)$ .

The kernel of the boundary operator is  $Z_n(X) = \text{ker}(\partial_n)$ , and is called the **group of singular** *n*-cycles. The image of the boundary operator is  $B_n(X) = \text{im}(\partial_{n+1})$ , and is called the **group of singular** *n*-boundaries.

It can also be shown that  $\partial_n \circ \partial_{n+1} = 0$ , implying  $B_n(X) \subseteq Z_n(X)$ . The *n*-th homology group of *X* is then defined as the factor group

$$H_n(X) = Z_n(X)/B_n(X).$$

The elements of  $H_n(X)$  are called **homology classes**.<sup>[2]</sup>

### Homotopy invariance

If *X* and *Y* are two topological spaces with the same homotopy type (i.e. are homotopy equivalent), then

$$H_n(X)\cong H_n(Y)$$

for all  $n \ge 0$ . This means homology groups are homotopy invariants, and therefore topological invariants.

In particular, if *X* is a connected <u>contractible space</u>, then all its homology groups are 0, except  $H_0(X) \cong \mathbb{Z}$ .

A proof for the homotopy invariance of singular homology groups can be sketched as follows. A continuous map  $f: X \rightarrow Y$  induces a homomorphism

$$f_{\sharp}: C_n(X) \to C_n(Y).$$

It can be verified immediately that

$$\partial f_{\sharp} = f_{\sharp} \partial,$$

i.e.  $f_{\#}$  is a chain map, which descends to homomorphisms on homology

 $f_*: H_n(X) \to H_n(Y).$ 

We now show that if *f* and *g* are homotopically equivalent, then  $f_* = g_*$ . From this follows that if *f* is a homotopy equivalence, then  $f_*$  is an isomorphism.

Let  $F : X \times [0, 1] \rightarrow Y$  be a homotopy that takes f to g. On the level of chains, define a homomorphism

 $P: C_n(X) \to C_{n+1}(Y)$ 

that, geometrically speaking, takes a basis element  $\sigma: \Delta^n \to X$  of  $C_n(X)$  to the "prism"  $P(\sigma): \Delta^n \times I \to Y$ . The boundary of  $P(\sigma)$  can be expressed as

$$\partial P(\sigma) = f_{\sharp}(\sigma) - g_{\sharp}(\sigma) - P(\partial \sigma).$$

So if  $\alpha$  in  $C_n(X)$  is an *n*-cycle, then  $f_{\#}(\alpha)$  and  $g_{\#}(\alpha)$  differ by a boundary:

$$f_{\sharp}(lpha) - g_{\sharp}(lpha) = \partial P(lpha),$$

i.e. they are homologous. This proves the claim. [3]

### Homology groups of common spaces

The table below shows the k-th homology groups  $H_k(X)$  of n-dimensional real projective spaces  $\mathbb{RP}^n$ , complex projective spaces,  $\mathbb{CP}^n$ , a point  $S^0$ , spheres  $S^n(n \ge 1)$ , and a 3-torus  $T^3$  with integer coefficients.

Space	Homotopy type	
<b>RP</b> <sup>n[4]</sup>	Z	k = 0 and $k = n$ odd
	$\mathbf{Z}/2\mathbf{Z}$	k odd, 0 < k < n
	0	otherwise
<b>CP</b> <sup><i>n</i>[5]</sup>	Z	k = 0,2,4,,2n
	0	otherwise
S <sup>0[6]</sup>	Z	k = 0
	0	otherwise
S <sup>n</sup>	Z	k = 0,n
	0	otherwise
T <sup>3[7]</sup>	Z	k = 0,3
	<b>Z</b> <sup>3</sup>	k = 1,2
	0	otherwise

### **Functoriality**

The construction above can be defined for any topological space, and is preserved by the action of continuous maps. This generality implies that singular homology theory can be recast in the language of <u>category theory</u>. In particular, the homology group can be understood to be a <u>functor</u> from the <u>category of topological spaces</u> **Top** to the <u>category of abelian groups</u> **Ab**.

Consider first that  $X \mapsto C_n(X)$  is a map from topological spaces to free abelian groups. This suggests that  $C_n(X)$  might be taken to be a functor, provided one can understand its action on the <u>morphisms</u> of **Top**. Now, the morphisms of **Top** are continuous functions, so if  $f : X \to Y$  is a continuous map of topological spaces, it can be extended to a homomorphism of groups

$$f_*:C_n(X) o C_n(Y)$$

by defining

$$f_*\left(\sum_i a_i\sigma_i
ight) = \sum_i a_i(f\circ\sigma_i)$$

where  $\sigma_i : \Delta^n \to X$  is a singular simplex, and  $\sum_i a_i \sigma_i$  is a singular *n*-chain, that is, an element of  $C_n(X)$ . This shows that  $C_n$  is a functor

 $C_n:\mathbf{Top}\to\mathbf{Ab}$ 

from the category of topological spaces to the category of abelian groups.

The boundary operator commutes with continuous maps, so that  $\partial_n f_* = f_* \partial_n$ . This allows the entire chain complex to be treated as a functor. In particular, this shows that the map  $X \mapsto H_n(X)$  is a functor

#### $H_n:\mathbf{Top}\to\mathbf{Ab}$

from the category of topological spaces to the category of abelian groups. By the homotopy axiom, one has that  $H_n$  is also a functor, called the homology functor, acting on **hTop**, the quotient homotopy category:

#### $H_n: \mathbf{hTop} \to \mathbf{Ab}.$

This distinguishes singular homology from other homology theories, wherein  $H_n$  is still a functor, but is not necessarily defined on all of **Top**. In some sense, singular homology is the "largest" homology theory, in that every homology theory on a <u>subcategory</u> of **Top** agrees with singular homology on that subcategory. On the other hand, the singular homology does not have the cleanest categorical properties; such a cleanup motivates the development of other homology theories such as <u>cellular</u> homology.

More generally, the homology functor is defined axiomatically, as a functor on an <u>abelian category</u>, or, alternately, as a functor on <u>chain complexes</u>, satisfying axioms that require a <u>boundary morphism</u> that turns <u>short</u> <u>exact sequences</u> into <u>long exact sequences</u>. In the case of singular homology, the homology functor may be factored into two pieces, a topological piece and an algebraic piece. The topological piece is given by

### $C_{\bullet}:\mathbf{Top}\to\mathbf{Comp}$

which maps topological spaces as  $X \mapsto (C_{\bullet}(X), \partial_{\bullet})$  and continuous functions as  $f \mapsto f_*$ . Here, then,  $C_{\bullet}$  is understood to be the singular chain functor, which maps topological spaces to the <u>category of chain complexes</u> **Comp** (or **Kom**). The category of chain complexes has chain complexes as its <u>objects</u>, and <u>chain maps</u> as its morphisms.

The second, algebraic part is the homology functor

### $H_n:\mathbf{Comp} o \mathbf{Ab}$

which maps

$$C_{ullet} \mapsto H_n(C_{ullet}) = Z_n(C_{ullet})/B_n(C_{ullet})$$

and takes chain maps to maps of abelian groups. It is this homology functor that may be defined axiomatically, so that it stands on its own as a functor on the category of chain complexes.

Homotopy maps re-enter the picture by defining homotopically equivalent chain maps. Thus, one may define the quotient category **hComp** or **K**, the homotopy category of chain complexes.

# **Coefficients in** *R*

Given any unital ring R, the set of singular n-simplices on a topological space can be taken to be the generators of a free R-module. That is, rather than performing the above constructions from the starting point of free abelian groups, one instead uses free R-modules in their place. All of the constructions go through with little or no change. The result of this is

### $H_n(X;R)$

which is now an <u>*R*-module</u>. Of course, it is usually *not* a free module. The usual homology group is regained by noting that

### $H_n(X;\mathbb{Z}) = H_n(X)$

when one takes the ring to be the ring of integers. The notation  $H_n(X; R)$  should not be confused with the nearly identical notation  $H_n(X, A)$ , which denotes the relative homology (below).

The <u>universal coefficient theorem</u> provides a mechanism to calculate the homology with R coefficients in terms of homology with usual integer coefficients using the short exact sequence

 $0 
ightarrow H_n(X;\mathbb{Z}) \otimes R 
ightarrow H_n(X;R) 
ightarrow Tor(H_{n-1}(X;\mathbb{Z}),R) 
ightarrow 0.$ 

where *Tor* is the <u>Tor functor</u>.<sup>[8]</sup> Of note, if *R* is torsion-free, then *Tor*(*G*, *R*) = 0 for any *G*, so the above short exact sequence reduces to an isomorphism between  $H_n(X;\mathbb{Z}) \otimes R$  and  $H_n(X;R)$ .

### **Relative homology**

For a subspace  $A \subset X$ , the <u>relative homology</u>  $H_n(X, A)$  is understood to be the homology of the quotient of the chain complexes, that is,

 $H_n(X,A) = H_n(C_{\bullet}(X)/C_{\bullet}(A))$ 

where the quotient of chain complexes is given by the short exact sequence

$$0 o C_{ullet}(A) o C_{ullet}(X) o C_{ullet}(X) / C_{ullet}(A) o 0.^{[9]}$$

### **Reduced homology**

The reduced homology of a space *X*, annotated as  $\tilde{H}_n(X)$  is a minor modification to the usual homology which simplifies expressions of some relationships and fulfils the intuiton that all homology groups of a point should be zero.

For the usual homology defined on a chain complex:

$$\cdots \xrightarrow{\partial_{n+1}} C_n \xrightarrow{\partial_n} C_{n-1} \xrightarrow{\partial_{n-1}} \cdots \xrightarrow{\partial_2} C_1 \xrightarrow{\partial_1} C_0 \xrightarrow{\partial_0} 0$$

To define the reduced homology, we augment the chain complex with an additional  $\mathbb{Z}$  between  $C_0$  and zero:

$$\cdots \xrightarrow{\partial_{n+1}} C_n \xrightarrow{\partial_n} C_{n-1} \xrightarrow{\partial_{n-1}} \cdots \xrightarrow{\partial_2} C_1 \xrightarrow{\partial_1} C_0 \xrightarrow{\epsilon} \mathbb{Z} \to 0$$

where  $\epsilon \left(\sum_{i} n_{i} \sigma_{i}\right) = \sum_{i} n_{i}$ . This can be justified by interpreting the empty set as "(-1)-simplex", which means that  $C_{-1} \simeq \mathbb{Z}$ .

The *reduced* homology groups are now defined by  $\tilde{H}_n(X) = \ker(\partial_n)/\operatorname{im}(\partial_{n+1})$  for positive *n* and  $\tilde{H}_0(X) = \ker(\epsilon)/\operatorname{im}(\partial_1)$ . [10]

For n > 0,  $H_n(X) = \tilde{H}_n(X)$ , while for n = 0,  $H_0(X) = \tilde{H}_0(X) \oplus \mathbb{Z}$ .

### Cohomology

By dualizing the homology <u>chain complex</u> (i.e. applying the functor Hom(-, *R*), *R* being any ring) we obtain a <u>cochain complex</u> with coboundary map  $\delta$ . The **cohomology groups** of *X* are defined as the homology groups of this complex; in a quip, "cohomology is the homology of the co [the dual complex]".

The cohomology groups have a richer, or at least more familiar, algebraic structure than the homology groups. Firstly, they form a differential graded algebra as follows:

- the graded set of groups form a graded R-module;
- this can be given the structure of a graded R-algebra using the cup product;
- the Bockstein homomorphism  $\beta$  gives a differential.

There are additional <u>cohomology</u> operations, and the cohomology algebra has addition structure mod p (as before, the mod p cohomology is the cohomology of the mod p cochain complex, not the mod p reduction of the cohomology), notably the <u>Steenrod algebra</u> structure.

# Betti homology and cohomology

Since the number of <u>homology theories</u> has become large (see <u>Category:Homology theory</u>), the terms **Betti** *homology* and **Betti cohomology** are sometimes applied (particularly by authors writing on <u>algebraic geometry</u>) to the singular theory, as giving rise to the <u>Betti numbers</u> of the most familiar spaces such as <u>simplicial complexes</u> and <u>closed manifolds</u>.

# Extraordinary homology

If one defines a homology theory axiomatically (via the <u>Eilenberg–Steenrod axioms</u>), and then relaxes one of the axioms (the *dimension axiom*), one obtains a generalized theory, called an <u>extraordinary homology theory</u>. These originally arose in the form of <u>extraordinary cohomology theories</u>, namely <u>K-theory</u> and <u>cobordism theory</u>. In this context, singular homology is referred to as **ordinary homology**.

### See also

- Derived category
- Excision theorem
- Hurewicz theorem
- Simplicial homology
- Cellular homology

# References

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