

**Postscript: why care about covering spaces?** The preceding discussion probably makes it clear that covering spaces play a central role in (combinatorial) group theory. It also plays a role in embedding problems; a common scenario is to have a map  $f : Y \rightarrow X$  which is injective on  $\pi_1$ , and we wish to know if we can lift  $f$  to a finite-sheeted covering so that the lifted map  $\tilde{f}$  is homotopic to an embedding. Information that is easier to obtain in the case of an embedding can then be passed down to gain information about the original map  $f$ . And covering spaces underlie the theory of analytic continuation in complex analysis; starting with a domain  $D \subseteq \mathbb{C}$ , what analytic continuation really builds is an (analytic) function from a covering space of  $D$  to  $\mathbb{C}$ . For example, the logarithm is really defined as a map from the universal cover of  $\mathbb{C} \setminus \{0\}$  to  $\mathbb{C}$ . The various “branches” of the logarithm refer to which sheet in this cover you are in.

**Homology theory:** Fundamental groups are a remarkably powerful tool for studying spaces; they capture a great deal of the global structure of a space, and so they are very good at distinguishing between homotopy-inequivalent spaces. In theory! But in practice, they suffer from the fact that deciding whether two groups are isomorphic or not is, in general, undecidable! Homology theory is designed to get around this deficiency; the theory, by design, builds (a sequence of) *abelian* groups  $H_i(X)$  from a topological space. And deciding whether or not two abelian groups, at least if you’re given a presentation for them, is, in the end, a matter of fairly routine linear algebra. Mostly because of the Fundamental Theorem of Finitely-generated Abelian groups; each such has a unique representation as  $\mathbb{Z}^m \oplus \mathbb{Z}_{m_1} \oplus \cdots \oplus \mathbb{Z}_{m_n}$  with  $m_{i+1} | m_i$  for every  $i$ .

There are also “higher” homotopy groups beyond the fundamental group  $\pi_1$ , (hence the name *pi-one*); elements are homotopy classes, rel boundary, of based maps  $(I^n, \partial I^n) \rightarrow (X, x_0)$ . Multiplication is again by concatenation. But unlike  $\pi_1$ , where we have a chance to compute it via Seifert-van Kampen, nobody, for example knows what all of the homotopy groups  $\pi_n(S^2)$  are (except that nearly all of them are non-trivial!). Like  $\pi_1$ , it describes, essentially, maps of  $S^n$  into  $X$  which don’t extend to maps of  $D^{n+1}$ , i.e., it turns the “ $n$ -dimensional holes” of  $X$  into a group.

Homology theory does the exact same thing, counting  $n$ -dimensional holes. In the end we will find it to be extremely computable; but it will require building a fair bit of machinery before it will become so transparent to calculate. But the short version is that the homology groups compute “cycles mod boundaries”, that is,  $n$ -dimensional objects/subsets that have no boundary (in the appropriate sense) modulo objects that are the boundary of  $(n+1)$ -dimensional ones. There are, in fact, probably as many ways to *define* homology groups as there are people actively working in the field; we will focus on two, simplicial homology and singular homology. The first is quick to define and compute, but hard to show is an invariant! The second is quick to see is an invariant, but, on the face of it, hard to compute! Luckily, for spaces where they are both defined, they are isomorphic. So, in the end, we get an invariant that is quick to compute. Of course, so is the invariant “4”; but this one will be a bit more informative than that....

First, simplicial homology. This is a sequence of groups defined for spaces for which they are easiest to define, which Hatcher calls  $\Delta$ -complexes. Basically, they are spaces defined by gluing simplices together using nice enough maps. More precisely, the *standard  $n$ -simplex*  $\Delta^n$  is the set of points  $\{(x_1, \dots, x_{n+1}) \in \mathbb{R}^{n+1} : \sum x_i = 1, x_i \geq 0 \text{ for all } i\}$ . This can also be expressed as convex linear combinations (literally, that’s the conditions on the  $x_i$ ’s) of the points  $e_i = (0, \dots, 0, 1, 0, \dots, 0)$ , the *vertices* of the standard simplex. More generally, an  $n$ -simplex is the set  $[v_0, \dots, v_n]$  of convex linear combinations of points  $v_0, \dots, v_n \in \mathbb{R}^k$  for which  $v_1 - v_0, \dots, v_n - v_0$  are linearly independent. Any bijection from the vertices of the standard simplex to the points  $v_0, \dots, v_n$  extends (linearly) to a homeomorphism of the simplices. The  $n+1$  *faces* of a simplex, each sitting opposite a vertex  $v_i$ , are obtained by setting the corresponding coefficient  $x_i$  to 0. Each forms an  $(n-1)$ -simplex, which we denote  $[v_0, \dots, v_{i-1}, v_{i+1}, \dots, v_n]$  or  $[v_0, \dots, \hat{v}_i, \dots, v_n]$ . A  $\Delta$ -complex  $X$  is a cell complex obtained by

gluing simplices together, but we insist on an extra condition: the restriction of the attaching map to any face is equal to a (lower-dimensional) cell. As before, we use the weak topology on the space; a set is open iff it's inverse image under the induced map of a cell into the complex is open. Each  $n$ -cell comes equipped with a (continuous) map  $\sigma : \Delta^n \rightarrow X$ , which is one-to-one on its interior, whose restriction to the boundary is the attaching map, and whose restriction to each face is the associated map for that  $(n-1)$ -simplex. We will typically blur the distinction between the map  $\sigma$  (called the *characteristic map* of the simplex) and its image, and denote the image by  $\sigma$  (or  $\sigma^n$ ), when this will cause no confusion, and call  $\sigma$  an  $n$ -simplex in  $X$ . When we feel the need for the distinction, we will use  $e^n$  for the image and  $\sigma^n$  for the map.

For example, taking our standard, identifications of the sides of a rectangle, cell structure for the 2-torus, and cutting the rectangle into two triangles (= 2-simplices) along a diagonal, we obtain a  $\Delta$ -structure with 2 2-simplices, 3 1-simplices, and 1 0-simplex. A genus  $g$  surface can be built, by cutting the  $2g$ -gon into triangles, with  $g+1$  2-simplices,  $3g$  1-simplices, and 1 0-simplex.

We typically think of building a  $\Delta$ -complex  $X$  inductively. The  $0$ -simplices (i.e., points), or *vertices*, form the 0-skeleton  $X^{(0)}$ .  $n$ -simplices  $\sigma^n = [v_0, \dots, v_n]$  attach to the  $(n-1)$ -skeleton to form the  $n$ -skeleton  $X^{(n)}$ ; the restriction of the attaching map to each face of  $\sigma^n$  is, by definition, an  $(n-1)$ -simplex in  $X$ . The attaching map is (by induction) really determined by a map  $\{v_0, \dots, v_n\} \rightarrow X^{(0)}$ , since this determines the attaching maps for the 1-simplices in the boundary of the  $n$ -simplex, which gives 1-simplices in  $X$ , which then give the attaching maps for the 2-simplices in the boundary, etc. Note that the reverse is not true; the vertices of two different  $n$ -simplices in  $X$  can be the same. For example, think of the 2-sphere as a pair of 2-simplices whose boundaries are glued by the identity.

The final detail that we need before defining (simplicial) homology groups is the notion of an *orientation* on a simplex of  $X$ . Each simplex  $\sigma^n$  is determined by a map  $f : \{v_0, \dots, v_n\} \rightarrow X^{(0)}$ ; an orientation on  $\sigma^n$  is an (equivalence class of) the ordered  $(n+1)$ -tuple  $(f(v_0), \dots, f(v_n)) = (V_0, \dots, V_n)$ . Another ordering of the same vertices represents the same orientation if there is an *even* permutation taking the entries of the first  $(n+1)$ -tuple to the second. This should be thought of as a generalization of the right-hand rule for  $\mathbb{R}^3$ , interpreted as orienting the vertices of a 3-simplex. Note that there are precisely two orientations on a simplex.

Now to define homology! We start by defining  $n$ -chains; these are (finite) formal linear combinations of the (oriented!)  $n$ -simplices of  $X$ , where  $-\sigma$  is interpreted as  $\sigma$  with the opposite (i.e., other) orientation. Adding formal linear combinations formally, we get the  $n$ -th *chain group*  $C_n(X) = \{\sum n_\alpha \sigma_\alpha : \sigma_\alpha \text{ an oriented } n\text{-simplex in } X\}$ . We next define a *boundary operator*  $\partial : C_n(X) \rightarrow C_{n-1}(X)$ , whose image will be the  $(n-1)$ -chains that are the “boundaries” of  $n$ -chains. We define it on the basis elements  $\sigma_\alpha = \sigma$  of  $C_n(X)$  as  $\partial\sigma = \sum (-1)^i \sigma|_{[v_0, \dots, \widehat{v_i}, \dots, v_n]}$ , where  $\sigma : [v_0, \dots, v_n] \rightarrow X$  is the characteristic map of  $\sigma_\alpha$ .  $\partial\sigma$  is therefore an alternating sum of the faces of  $\sigma$ . The point that really make this definition go is that we need *oriented* simplices, so that we know what the  $i$ -th face of  $\sigma$  is (the one opposite the  $i$ -th vertex). We then extend the definition by linearity to all of  $C_n(X)$ . When a notation indicating dimension is needed, we write  $\partial = \partial_n$ .

This definition is cooked up to make the maxim “boundaries have no boundary” true; that is  $\delta_{n-1} \circ \delta_n = 0$ , the 0 map. This is because, for any simplex  $\sigma = [v_0, \dots, v_n]$ ,

$$\begin{aligned} \delta \circ \delta(\sigma) &= \delta\left(\sum_{i=0}^n (-1)^i \sigma|_{[v_0, \dots, \widehat{v_i}, \dots, v_n]}\right) \\ &= \left(\sum_{j < i} (-1)^j (-1)^i \sigma|_{[v_0, \dots, \widehat{v_j}, \dots, \widehat{v_i}, \dots, v_n]}\right) + \left(\sum_{j > i} (-1)^{j-1} (-1)^i \sigma|_{[v_0, \dots, \widehat{v_i}, \dots, \widehat{v_j}, \dots, v_n]}\right) \end{aligned}$$

The distinction between the two pieces is that in the second part,  $v_j$  is actually the  $(j-1)$ -st vertex of the face. Switching the roles of  $i$  and  $j$  in the second sum, we find that the two are negatives of one another, so they sum to 0, as desired.

And this little calculation is all that it takes to define homology groups! What this tells us is that  $\text{im}(\delta_{n+1}) \subseteq \ker(\delta_n$  for every  $n$ .  $\ker(\delta_n = Z_n(X)$  are called the *n-cycles* of  $X$ ; they are the  $n$ -chains with 0 (i.e., empty) boundary. They form a (free) abelian subgroup of  $C_n(X)$ .  $\text{im}(\delta_{n+1} = B_n(X)$  are the *n-boundaries* of  $X$ ; they are, of course, the boundaries of  $(n+1)$ -chains in  $X$ . The  $n$ -th homology group of  $X$ ,  $H_n(X)$  is the quotient  $Z_n(X)/B_n(X)$  ; it is an abelian group.